

Accession	Sequence	Position
Oy	AGTATTAAATAGTAAATCAATTACGGGGTCAATTAGTCATAGCCCAATATATGAGATTTCCG	61
Dd	AGTATTAAATAGTAAATCAATTACGGGGTCAATTAGTCATAGCCCAATATATGAGATTTCCG	893
Oy	GTTACATAACTTACGGTAAATTGCGCCCGCGGTGACGCCCAAGACGCCCGCCCAATG	121
Dd	GTTACATAACTTACGGTAAATGCGCCCGCGGTGACGCCCAAGACGCCCGCCCAATG	953
Oy	ACGTCATAATAGAGCTATGTTCCCATATGTAAGCCCAATAGGACCTTTCATATGAGCTAA	181
Dd	ACGTCATAATAGAGCTATGTTCCCATATGTAAGCCCAATAGGACCTTTCATATGAGCTAA	1013
Oy	TGGGTGAGATTTACGGTAACAGCCCA-TTGGCAGTACATCAAGTATCATATATGCCA	240
Dd	TGGGTGAGATTTACGGTAACAGCCCA-TTGGCAGTACATCAAGTATCATATATGCCA	1014

OY	241	AGAGGAGGCCCCCTATTGACGCAATGAGCGGTAATGG-----	276
Db	1074	AGTACGCCCCCTATTGACGCTCAATGACGGTAATAGCCCGCCTGGCATTATGCCCCAGTAC	1133
OY	277	-----	276
Db	1134	ATGACCTTATGGAGCTTTCCTACTTGGCAGTACATCTAGCTATTATGATCATCGCTATTAC	1193
OY	277	-----	276
Db	1194	ATGCATGCTGAGGTGAGCCCCACAGCTTCGCTCTCACTCTCCCATCTCCCCCCTCCCC	1255
OY	277	-----ATGCAGTATTTTGTGACGCGATGGGGG---GG	306
Db	1254	ACCCCCAATTTTGTATTATTATTATTATTATTATTATTGTCAGCGCATGCGGGGCGGGGG	1313
OY	307	GGGGGGGGGGGGCGCCCGCAGCGGGGGGGGGGGCGAGGGGGGGGGGGGGGGCGACGCG	366
Db	1314	GGGGGGGGGGGGCGCCCGCAGCGGGGGGGGGGGCGAGGGGGGGGGGGGGGGCGACGCG	1373
OY	367	GAGAGGTGCGGCGGACAGCAATCAGAGCGGCGGCTGCCGAAAGTTCTTTTATGGCGAG	426
Db	1374	GAGAGGTGCGGCGGACAGCAATCAGAGCGGCGGCTGCCGAAAGTTCTTTTATGGCGAG	1433
OY	427	GGGGGGGGGGGGCGGCGCCCTATTAAAAAGCAACGCGGCGGGGGGGAGTGGCTGCGCG	486
Db	1434	GCGGGGGGGGGGGCGGCGCCCTATTAAAAAGCAACGCGGCGGGGGGGAGTGGCTGCGCG	1493
OY	487	CTGACCTTGGCCCGCTGCGCCGCGCTCCGCGCGGCGCTCGCGCGCGCGCCCGCGGCTCGACT	546
Db	1494	CTGACCTTGGCCCGCTGCGCCGCGCTCCGCGCGGCGCTCGCGCGCGCGCCCGCGGCTCGACT	1555
OY	547	GACGCGCGT	554
Db	1554	GACGCGCGT	1561

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Db	349	ACGTCATTAATGAGCTAATGTTCCCATAGTAAGCCCAATAGGAGCTTCCATTCAGCTCAA	408
0Y	182	TGGTGTCAGTATTTCACGTAACACTGCCCC-TTGGCACTACATCAAGTGTATCATATGCCA	240
Db	409	TGGGTGGAGTATTTCACGTAACACTGCCCCACTGGCACTACATCAAGTGTATCATATGCCA	468
0Y	241	AGTACGCCCCCTATTGACGTCAATGACGGTAAATGG	276
Db	469	AGTACGCCCCCTATTGACGTCAATGACGGTAAATGG	504

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    RESULT 3
    US-09-897-006-4
    ; Sequence 4, Application US/09897006
    ; Patent No. US20020106729A1
    ; GENERAL INFORMATION:
    ; APPLICANT: Black, Gregory
    ; TITLE OF INVENTION: Expression Vectors
    ; FILE REFERENCE: GALA-06415
    ; CURRENT APPLICATION NUMBER: US/09/897,006
    ; CURRENT FILING DATE: 2001-06-29
    ; PRIOR APPLICATION NUMBER: 60/215,851
    ; PRIOR FILING DATE: 2000-07-03
    ; NUMBER OF SEQ ID NOS: 36
    ; SOFTWARE: PatentIn version 3.0
    ; SEQ ID NO 4
    ; LENGTH: 4207
    ; TYPE: DNA
    ; ORGANISM: Artificial Sequence
    ; FEATURE:
    ; OTHER INFORMATION: Synthetic
    US-09-897-006-4
  
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Query Match	Similarity	96.1%	Score 256	DB 10	Length 4207
Best Local	Similarity 97.8%	Pred. No. 1,4e-49			
Matches 270	Conservative 0	Mismatches 5	Indels 1	Gaps 1	
Qy	2	ACTATTATTAAGTAAATCAATTAACGGGGGTCATTAGTTCATATGCCCATTATGGAGTTCCG	61		
Db	154	ACTATTATTAAGTAAATCAATTAACGGGGGTCATTAGTTCATATGCCCATTATGGAGTTCCG	213		
Qy	62	GTTACATACCTACAGGTAATATGGCCCGCGGCTGACGCCGCCACAGACCCCGCCCATTTG	121		
Db	214	GTTACATACCTACAGGTAATATGGCCCGCGGCTGACGCCGCCACAGACCCCGCCCATTTG	273		
Qy	122	ACGTCATATATGACGTATGTTCCCATAGTAACGCCCATATGGACATTTCCATTGAGCTCA	181		
Db	274	ACGTCATATATGACGTATGTTCCCATAGTAACGCCCATATGGACATTTCCATTGAGCTCA	333		
Qy	182	TGGGTGGCAATATTTACGGTAATGTCGCCA-TTGGCAGTACATCAATGATATATATATGCA	240		
Db	334	TGGGTGGCAATATTTACGGTAATGTCGCCA-TTGGCAGTACATCAATGATATATATATGCA	393		
Qy	241	AGTAGCCCCCTATTGACGTCAATGACGGTAATGG	276		
Db	394	AGTAGCCCCCTATTGACGTCAATGACGGTAATGG	429		

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, RESULT 4
, US-09-897-006-5
, Sequence 5, Application US/09897006
, Patent No. US20020106729A1
, GENERAL INFORMATION:
, APPLICANT: Black, Gregory
, TITLE OF INVENTION: Expression Vectors
, FILE REFERENCE: GALA-06415
, CURRENT APPLICATION NUMBER: US/09/897, 006
, CURRENT FILING DATE: 2001-06-29
, PRIOR APPLICATION NUMBER: 60/215,851
, PRIOR FILING DATE: 2000-07-03
, NUMBER OF SEQ ID NOS: 36
, SOFTWARE: PatentIn version 3.0

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SEQ ID NO 5
LENGTH: 4210
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Synthetic
US-09-897-006-5

Query Match 46.1%; Score 256; DB 10; Length 4210;
Best Local Similarity 97.8%; Pred. No. 1.4e-49;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 2 AGTTTATATAGTAATCAATTAACGGGGGTCATTAAGTTCATAGCCCATATATGAGTTCCGC 61
DB 153 AGTTTATATAGTAATCAATTAACGGGGGTCATTAAGTTCATAGCCCATATATGAGTTCCGC 212
QY 62 GTTACATTAACCTTAAGGTAATGCGCCGCGCTGACGCCCAAGACGCCCGCCCATTTG 121
DB 213 GTTACATTAACCTTAAGGTAATGCGCCGCGCTGACGCCCAAGACGCCCGCCCATTTG 272
QY 122 ACCTCAATATAGCAATGATGTCCTCCATAGTACGCAATGAGGACTTCCATTAAGCTCAA 181
DB 273 ACCTCAATATAGCAATGATGTCCTCCATAGTACGCAATGAGGACTTCCATTAAGCTCAA 332
QY 182 TGGGTGAGTATTTACGCTAACTGCCCA-TTGGCAGTACATCAATGATATATATGCCA 240
DB 333 TGGGTGAGTATTTACGCTAACTGCCCACTTGGCAGTACATCAATGATATATATGCCA 392
QY 241 AGTACGCCCCCTATTGACGTCAATGACGGTAATGG 276
DB 393 AGTACGCCCCCTATTGACGTCAATGACGGTAATGG 428

RESULT 5

US-09-759-960-7
Sequence 7, Application US/09759960
Patent No. US2001000639A1

GENERAL INFORMATION:

APPLICANT: Urban, Robert G.
APPLICANT: Chicz, Roman M.
APPLICANT: Collins, Edward J.
APPLICANT: Hedley, Mary Lynn
TITLE OF INVENTION: IMMUNOGENIC PEPTIDES FROM THE HPV E7
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 225 Franklin Street
CITY: Boston
STATE: MA
COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/759,960
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/169,425
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 08191/004002
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-5070
TELEFAX: 617-543-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:

LENGTH: 4665 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: DNA
US-09-759-960-7

Query Match 46.1%; Score 256; DB 10; Length 4665;
Best Local Similarity 97.8%; Pred. No. 1.4e-49;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 2 AGTTATTAATAGTAATCAATTAACGGGGTCAATTAAGTTCATAGCCCATATATGAGTTCCGC 61
DB 2625 AGTTATTAATAGTAATCAATTAACGGGGTCAATTAAGTTCATAGCCCATATATGAGTTCCGC 2684
QY 62 GTTACATTAACCTTAAGGTAATGCGCCGCGCTGACGCCCAAGACGCCCGCCCATTTG 121
DB 2685 GTTACATTAACCTTAAGGTAATGCGCCGCGCTGACGCCCAAGACGCCCGCCCATTTG 2744
QY 122 ACCTCAATATAGCAATGATGTCCTCCATAGTACGCAATGAGGACTTCCATTAAGCTCAA 181
DB 2745 ACCTCAATATAGCAATGATGTCCTCCATAGTACGCAATGAGGACTTCCATTAAGCTCAA 2804
QY 182 TGGGTGAGTATTTACGCTAACTGCCCA-TTGGCAGTACATCAATGATATATATGCCA 240
DB 2805 TGGGTGAGTATTTACGCTAACTGCCCACTTGGCAGTACATCAATGATATATATGCCA 2864
QY 241 AGTACGCCCCCTATTGACGTCAATGACGGTAATGG 276
DB 2865 AGTACGCCCCCTATTGACGTCAATGACGGTAATGG 2900

RESULT 6

US-09-812-133-5
Sequence 5, Application US/09812133
Patent No. US20020065240A1

GENERAL INFORMATION:

APPLICANT: Thomas, Kenneth A., Jr.
APPLICANT: Kendall, Richard L.
APPLICANT: Bett, Andrew J.
APPLICANT: Huckle, William R.
TITLE OF INVENTION: GENE THERAPY FOR STIMULATION OF
FILE OF INVENTION: ANGIOGENESIS
FILE REFERENCE: 20073P
CURRENT APPLICATION NUMBER: US/09/812,133
CURRENT FILING DATE: 2001-03-19
PRIOR APPLICATION NUMBER: PCT/US98/22668
PRIOR FILING DATE: 1998-10-23
PRIOR APPLICATION NUMBER: 60/063,629
PRIOR FILING DATE: 1997-10-27
NUMBER OF SEQ ID NOS: 5
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 5
LENGTH: 4864
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: E. coli
US-09-812-133-5

Query Match 46.1%; Score 256; DB 10; Length 4864;
Best Local Similarity 97.8%; Pred. No. 1.4e-49;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 2 AGTTATTAATAGTAATCAATTAACGGGGTCAATTAAGTTCATAGCCCATATATGAGTTCCGC 61
DB 338 AGTTATTAATAGTAATCAATTAACGGGGTCAATTAAGTTCATAGCCCATATATGAGTTCCGC 397
QY 62 GTTACATTAACCTTAAGGTAATGCGCCGCGCTGACGCCCAAGACGCCCGCCCATTTG 121
DB 398 GTTACATTAACCTTAAGGTAATGCGCCGCGCTGACGCCCAAGACGCCCGCCCATTTG 457
QY 122 ACCTCAATATAGCAATGATGTCCTCCATAGTACGCAATGAGGACTTCCATTAAGCTCAA 181

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Db 458 ACGCAATATGAGCTATGTTCCCATGTACGCCCATAGGAGCTTTCATTCAGCTCAA 517
Qy 182 TGGGTGAGATTTATTCAGTAACAGCCCA-TTGGCAGTACATCAAGTGTATCATATGCCA 240
|||||
Db 518 TGGGTGAGATTTATTCAGTAACAGCCCACTTGGCAGTACATCAAGTGTATCATATGCCA 577
Qy 241 AGTACGCCCTATTCAGCTCAATGAGGTAATGG 276
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Db 578 AGTACGCCCTATTCAGCTCAATGAGGTAATGG 613
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RESULT 7

US-09-966-976A-5
; Sequence 5, Application US/09966976A
; Patent No. US20020168649A1
; GENERAL INFORMATION:
; APPLICANT: Ferrick, David A.
; APPLICANT: Swift, Susan E.
; APPLICANT: Armstrong, Randall
; APPLICANT: Fox, Bryan
; TITLE OF INVENTION: Secretion and Compositions for Screening for Modulators and Ige Syt
; FILE REFERENCE: A-66038-4/RMS/JUD/DLR
; CURRENT FILING DATE: 2001-09-27
; PRIOR FILING DATE: 1998-05-12
; NUMBER OF SEQ ID NOS: 19
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
; LENGTH: 5713
; TYPE: DNA
; ORGANISM: Artificial sequence
; FEATURE:
; OTHER INFORMATION: synthetic
US-09-966-976A-5

Query Match 46.1%; Score 256; DB 9; Length 5713;
Best Local Similarity 97.8%; Pred. No. 1.4e-49;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

Qy 2 AGTATTTATATGATCAATATACGGGGTCATTAGTTCATACCCCATATATGAGTTCCGC 61
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Db 102 AGTATTTATATGATCAATATACGGGGTCATTAGTTCATACCCCATATATGAGTTCCGC 161
Qy 62 GTTACATACCTAGCTGAATATGGCCCGCGGTGACCGCCCAAGACCCCGCCCATGG 121
|||||
Db 162 GTTACATACCTAGCTGAATATGGCCCGCGGTGACCGCCCAAGACCCCGCCCATGG 221
Qy 122 ACGTCAATATGAGCTATGTTCCCATAGTAAAGCCCAATAGGAGACTTTCATTCAGCTCAA 181
|||||
Db 222 ACGTCAATATGAGCTATGTTCCCATAGTAAAGCCCAATAGGAGACTTTCATTCAGCTCAA 281
Qy 182 TGGGTGAGATTTATTCAGTAACAGCCCA-TTGGCAGTACATCAAGTGTATCATATGCCA 240
|||||
Db 282 TGGGTGAGATTTATTCAGTAACAGCCCACTTGGCAGTACATCAAGTGTATCATATGCCA 341
Qy 241 AGTACGCCCTATTCAGCTCAATGAGGTAATGG 276
|||||
Db 342 AGTACGCCCTATTCAGCTCAATGAGGTAATGG 377
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RESULT 8

US-09-963-206B-5
; Sequence 5, Application US/09963206B
; Patent No. US20020123076A1
; GENERAL INFORMATION:
; APPLICANT: Ferrick, David A.
; APPLICANT: Swift, Susan E.
; APPLICANT: Armstrong, Randall
; APPLICANT: Fox, Bryan
; TITLE OF INVENTION: Methods and Compositions for Screening for Modulators and Ige Syt

|||||
; TITLE OF INVENTION: Secretion and Switch Rearrangement
; FILE REFERENCE: A-66038-3/RMS/JUD/DLR
; CURRENT APPLICATION NUMBER: US/09/963, 206B
; CURRENT FILING DATE: 2001-09-25
; PRIOR APPLICATION NUMBER: US 09/076,624
; PRIOR FILING DATE: 1998-05-12
; NUMBER OF SEQ ID NOS: 19
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
; LENGTH: 5713
; TYPE: DNA
; ORGANISM: Artificial sequence
; FEATURE:
; OTHER INFORMATION: synthetic
US-09-963-206B-5

Query Match 46.1%; Score 256; DB 10; Length 5713;
Best Local Similarity 97.8%; Pred. No. 1.4e-49;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

Qy 2 AGTATTTATATGATCAATATACGGGGTCATTAGTTCATAGCCCATATATGAGTTCCGC 61
|||||
Db 102 AGTATTTATATGATCAATATACGGGGTCATTAGTTCATAGCCCATATATGAGTTCCGC 161
Qy 62 GTTACATACCTAGCTGAATATGGCCCGCGGTGACCGCCCAAGACCCCGCCCATGG 121
|||||
Db 162 GTTACATACCTAGCTGAATATGGCCCGCGGTGACCGCCCAAGACCCCGCCCATGG 221
Qy 122 ACGTCAATATGAGCTATGTTCCCATAGTAAAGCCCAATAGGAGACTTTCATTCAGCTCAA 181
|||||
Db 222 ACGTCAATATGAGCTATGTTCCCATAGTAAAGCCCAATAGGAGACTTTCATTCAGCTCAA 281
Qy 182 TGGGTGAGATTTATTCAGTAACAGCCCA-TTGGCAGTACATCAAGTGTATCATATGCCA 240
|||||
Db 282 TGGGTGAGATTTATTCAGTAACAGCCCACTTGGCAGTACATCAAGTGTATCATATGCCA 341
Qy 241 AGTACGCCCTATTCAGCTCAATGAGGTAATGG 276
|||||
Db 342 AGTACGCCCTATTCAGCTCAATGAGGTAATGG 377
|||||

RESULT 9

US-09-966-976A-4
; Sequence 4, Application US/09966976A
; Patent No. US20020168649A1
; GENERAL INFORMATION:
; APPLICANT: Ferrick, David A.
; APPLICANT: Swift, Susan E.
; APPLICANT: Armstrong, Randall
; APPLICANT: Fox, Bryan
; TITLE OF INVENTION: Methods and Compositions for Screening for Modulators and Ige
; FILE REFERENCE: A-66038-4/RMS/JUD/DLR
; CURRENT FILING DATE: 2001-09-27
; PRIOR FILING DATE: 1998-05-12
; NUMBER OF SEQ ID NOS: 19
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 4
; LENGTH: 6219
; TYPE: DNA
; ORGANISM: Artificial sequence
; FEATURE:
; OTHER INFORMATION: synthetic
US-09-966-976A-4

Query Match 46.1%; Score 256; DB 9; Length 6219;
Best Local Similarity 97.8%; Pred. No. 1.5e-49;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

Qy 2 AGTATTTATATGATCAATATACGGGGTCATTAGTTCATAGCCCATATATGAGTTCCGC 61
|||||

Db 102 AGTATTATATAGTATCAATTAACGGGGTCAATTAGTTCATACGCCCATATATATGAGTTCCGC 161
 Oy 62 GTTACATTAATTAAGGTAATTTGGCCCGCTGACCGCCCAACGACGCCGCCCATATG 121
 Db 162 GTTACATTAATTAAGGTAATTTGGCCCGCTGACCGCCCAACGACGCCGCCCATATG 221
 Oy 122 AGCTCAATTAATGACGATGTTCCCATATGTAACGCCCAATAGGACCTTCCATTGACGTCAA 181
 Db 222 AGCTCAATTAATGACGATGTTCCCATATGTAACGCCCAATAGGACCTTCCATTGACGTCAA 281
 Oy 182 TGGGTGAGTATTTACGCTAAATGCGCCA-TTGGCAGTACATCAAGTATATATATGCA 240
 Db 282 TGGGTGAGTATTTACGCTAAATGCGCCA-TTGGCAGTACATCAAGTATATATATGCA 341
 Oy 241 AGTACGCCGCCCTATTTAGCTCAATGAGCGTAAATGG 276
 Db 342 AGTACGCCGCCCTATTTAGCTCAATGAGCGTAAATGG 377

RESULT 10
 US-09-963-2068-4
 : Sequence 4, Application US/099632068
 : Patent No. US20020123076A1
 : GENERAL INFORMATION:
 : APPLICANT: Ferrick, David A.
 : APPLICANT: Swift, Susan E.
 : APPLICANT: Armstrong, Randall
 : APPLICANT: Fox, Bryan
 : TITLE OF INVENTION: Methods and Compositions for Screening for Modulators and Ige Syn
 : FILE REFERENCE: A-66038-3/RMS/JJD/DLR
 : CURRENT APPLICATION NUMBER: US/09/963, 2068
 : PRIOR FILING DATE: 2001-09-25
 : PRIOR APPLICATION NUMBER: US 09/076, 624
 : NUMBER OF SEQ ID NOS: 19
 : SOFTWARE: PatentIn version 3.1
 : SEQ ID NO 4
 : LENGTH: 6219
 : TYPE: DNA
 : ORGANISM: Artificial sequence
 : FEATURE:
 : OTHER INFORMATION: synthetic
 : US-09-963-2068-4

Query Match 46.1%; Score 256; DB 10; Length 6219;
 Best Local Similarity 97.8%; Pred. No. 1.5e-49;
 Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

Oy 2 AGTATTATATAGTATCAATTAACGGGGTCAATTAGTTCATACGCCCATATATGAGTTCCGC 61
 Db 102 AGTATTATATAGTATCAATTAACGGGGTCAATTAGTTCATACGCCCATATATGAGTTCCGC 161
 Oy 62 GTTACATTAATTAAGGTAATTTGGCCCGCTGACCGCCCAACGACGCCGCCCATATG 121
 Db 162 GTTACATTAATTAAGGTAATTTGGCCCGCTGACCGCCCAACGACGCCGCCCATATG 221
 Oy 122 AGCTCAATTAATGACGATGTTCCCATATGTAACGCCCAATAGGACCTTCCATTGACGTCAA 181
 Db 222 AGCTCAATTAATGACGATGTTCCCATATGTAACGCCCAATAGGACCTTCCATTGACGTCAA 281
 Oy 182 TGGGTGAGTATTTACGCTAAATGCGCCA-TTGGCAGTACATCAAGTATATATATGCA 240
 Db 282 TGGGTGAGTATTTACGCTAAATGCGCCA-TTGGCAGTACATCAAGTATATATATGCA 341
 Oy 241 AGTACGCCGCCCTATTTAGCTCAATGAGCGTAAATGG 276
 Db 342 AGTACGCCGCCCTATTTAGCTCAATGAGCGTAAATGG 377

RESULT 11
 US-09-897-006-13
 : Sequence 13, Application US/09897006

Patent No. US20020106729A1
 : GENERAL INFORMATION:
 : APPLICANT: Bleck, Gregory
 : TITLE OF INVENTION: Expression Vectors
 : FILE REFERENCE: GALA-06415
 : CURRENT APPLICATION NUMBER: US/09/897, 006
 : PRIOR FILING DATE: 2001-06-29
 : PRIOR APPLICATION NUMBER: 60/215, 851
 : NUMBER OF SEQ ID NOS: 36
 : SOFTWARE: PatentIn version 3.0
 : SEQ ID NO 13
 : LENGTH: 6255
 : TYPE: DNA
 : ORGANISM: Artificial Sequence
 : FEATURE:
 : OTHER INFORMATION: Synthetic
 : US-09-897-006-13

Query Match 46.1%; Score 256; DB 10; Length 6255;
 Best Local Similarity 97.8%; Pred. No. 1.5e-49;
 Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

Oy 2 AGTATTATATAGTATCAATTAACGGGGTCAATTAGTTCATACGCCCATATATGAGTTCCGC 61
 Db 2805 AGTATTATATAGTATCAATTAACGGGGTCAATTAGTTCATACGCCCATATATGAGTTCCGC 2864
 Oy 62 GTTACATTAATTAAGGTAATTTGGCCCGCTGACCGCCCAACGACGCCGCCCATATG 121
 Db 2865 GTTACATTAATTAAGGTAATTTGGCCCGCTGACCGCCCAACGACGCCGCCCATATG 2924
 Oy 122 AGCTCAATTAATGACGATGTTCCCATATGTAACGCCCAATAGGACCTTCCATTGACGTCAA 181
 Db 2925 AGCTCAATTAATGACGATGTTCCCATATGTAACGCCCAATAGGACCTTCCATTGACGTCAA 2984
 Oy 182 TGGGTGAGTATTTACGCTAAATGCGCCA-TTGGCAGTACATCAAGTATATATATGCA 240
 Db 2985 TGGGTGAGTATTTACGCTAAATGCGCCA-TTGGCAGTACATCAAGTATATATATGCA 3044
 Oy 241 AGTACGCCGCCCTATTTAGCTCAATGAGCGTAAATGG 276
 Db 3045 AGTACGCCGCCCTATTTAGCTCAATGAGCGTAAATGG 3080

RESULT 12
 US-09-982-610-17
 : Sequence 17, Application US/09982610
 : Patent No. US20020146420A1
 : GENERAL INFORMATION:
 : APPLICANT: Genentech, Inc.
 : APPLICANT: Bennett, Brian D.
 : APPLICANT: Goeddel, David
 : APPLICANT: Lee, James M.
 : APPLICANT: Mathews, William
 : APPLICANT: Tsai, Siao Ping
 : APPLICANT: Wood, William I.

TITLE OF INVENTION: PROTEIN TYROSINE KINASE AGONIST ANTIBODIES
 NUMBER OF SEQUENCES: 45
 CORRESPONDENCE ADDRESS:
 ADDRESS: Genentech, Inc.
 STREET: 460 Point San Bruno Blvd
 CITY: South San Francisco
 STATE: California
 COUNTRY: USA
 ZIP: 94080
 COMPUTER READABLE FORM:
 MEDIUM TYPE: 3.5 Inch, 1.44 MB floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Winpatin (Genentech)
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/982, 610
 FILING DATE: 17-Oct-2001

CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/446,648

FILING DATE: 1996-MAY-23

APPLICATION NUMBER: 08/222616

FILING DATE: 04-APR-1994

ATTORNEY/AGENT INFORMATION:

NAME: Lee, Wendy M.

REGISTRATION NUMBER: 40,378

REFERENCE/DOCKET NUMBER: P0821P3PCT

TELECOMMUNICATION INFORMATION:

TELEPHONE: 415/225-1994

TELEFAX: 415/952-9881

TELEX: 910/371-7168

INFORMATION FOR SEQ ID NO: 17:

SEQUENCE CHARACTERISTICS:

LENGTH: 6827 base pairs

TYPE: Nucleic Acid

STRANDEDNESS: Single

TOPOLOGY: Linear

SEQUENCE DESCRIPTION: SEQ ID NO: 17:

US-09-982-610-17

Query Match 46.1%; Score 256; DB 10; Length 6827;

Best Local Similarity 97.8%; Pred. No. 1.5e-49;

Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 2 AGTTATTAATGATATCATATACGGGGTCAATGATGATACCCATATATGAGTTCCGC 61

DB 31 AGTTATTAATGATATCATATACGGGGTCAATGATGATACCCATATATGAGTTCCGC 90

QY 62 GTTACATACTTACGTAATATGCGCGCGGCTGACGCCACGACCCCGCCCATTTG 121

DB 91 GTTACATACTTACGTAATATGCGCGCGGCTGACGCCACGACCCCGCCCATTTG 150

QY 122 ACGTCAATATGACGTAATGTTCCCATATGACGTAATGAGGACTTTCATGAGCTCA 181

DB 151 ACGTCAATATGACGTAATGTTCCCATATGACGTAATGAGGACTTTCATGAGCTCA 210

QY 182 TGGGTGAGTATTTACGTAATGCGCGCGGCTGACGCCACGACCCCGCCCATTTG 240

DB 211 TGGGTGAGTATTTACGTAATGCGCGCGGCTGACGCCACGACCCCGCCCATTTG 270

QY 241 AGRACGCCCCCTATTGACGTCAATGACGTAATGCG 276

DB 271 AGRACGCCCCCTATTGACGTCAATGACGTAATGCG 306

RESULT 13

US-09-872-733-15

Sequence 15, Application US/09872733

Patent No. US20010036655A1

GENERAL INFORMATION:

APPLICANT: The Government of the United States of America, as

TITLE OF INVENTION: MOLECULAR CLONES WITH MODIFIED HIV GAG/POL, SIV GAG AND

FILE REFERENCE: 2026-4287051 HIV GAG/POL, SIV GAG & ENV

CURRENT FILING DATE: 2001-06-01

PRIOR APPLICATION NUMBER: US/09/872,733

PRIOR FILING DATE: 2000-12-22

PRIOR APPLICATION NUMBER: PCT/US00/34985

PRIOR FILING DATE: 1999-12-23

NUMBER OF SEQ ID NOS: 19

SOFTWARE: PatentIn Ver. 2.1

SEQ ID NO 15

LENGTH: 6978

TYPE: DNA

ORGANISM: Artificial Sequence

FEATURE:

OTHER INFORMATION: Description of Artificial Sequence: DNA sequence

US-09-872-733-15

Query Match 46.1%; Score 256; DB 9; Length 6978;

Best Local Similarity 97.8%; Pred. No. 1.5e-49;

Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 2 AGTTATTAATGATATCATATACGGGGTCAATGATGATACCCATATATGAGTTCCGC 61

DB 96 AGTTATTAATGATATCATATACGGGGTCAATGATGATACCCATATATGAGTTCCGC 155

QY 62 GTTACATACTTACGTAATATGCGCGCGGCTGACGCCACGACCCCGCCCATTTG 121

DB 156 GTTACATACTTACGTAATATGCGCGCGGCTGACGCCACGACCCCGCCCATTTG 215

QY 122 ACGTCAATATGACGTAATGTTCCCATATGACGTAATGAGGACTTTCATGAGCTCA 181

DB 216 ACGTCAATATGACGTAATGTTCCCATATGACGTAATGAGGACTTTCATGAGCTCA 275

QY 182 TGGGTGAGTATTTACGTAATGCGCGCGGCTGACGCCACGACCCCGCCCATTTG 240

DB 276 TGGGTGAGTATTTACGTAATGCGCGCGGCTGACGCCACGACCCCGCCCATTTG 335

QY 241 AGRACGCCCCCTATTGACGTCAATGACGTAATGCG 276

DB 336 AGRACGCCCCCTATTGACGTCAATGACGTAATGCG 371

RESULT 14

US-10-001-189-45

Sequence 45, Application US/10001189

Patent No. US20020173634A1

GENERAL INFORMATION:

APPLICANT: FRASER JR., MALCOLM J.

APPLICANT: LI, XU

TITLE OF INVENTION: BEAM, TERESA

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TRANSPOSITION USING

FILE REFERENCE: 835910-92098

CURRENT APPLICATION NUMBER: US/10/001,189

PRIOR FILING DATE: 2001-10-30

PRIOR APPLICATION NUMBER: 60/244,984

PRIOR FILING DATE: 2000-11-01

PRIOR APPLICATION NUMBER: 60/244,677

PRIOR FILING DATE: 2000-10-31

NUMBER OF SEQ ID NOS: 70

SOFTWARE: PatentIn Ver. 2.1

SEQ ID NO 45

LENGTH: 6984

TYPE: DNA

ORGANISM: Artificial Sequence

FEATURE:

OTHER INFORMATION: Description of Artificial Sequence:

US-10-001-189-45

Query Match 46.1%; Score 256; DB 9; Length 6984;

Best Local Similarity 97.8%; Pred. No. 1.5e-49;

Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 2 AGTTATTAATGATATCATATACGGGGTCAATGATGATACCCATATATGAGTTCCGC 61

DB 2 AGTTATTAATGATATCATATACGGGGTCAATGATGATACCCATATATGAGTTCCGC 61

QY 62 GTTACATACTTACGTAATATGCGCGCGGCTGACGCCACGACCCCGCCCATTTG 121

DB 62 GTTACATACTTACGTAATATGCGCGCGGCTGACGCCACGACCCCGCCCATTTG 121

QY 122 ACGTCAATATGACGTAATGTTCCCATATGACGTAATGAGGACTTTCATGAGCTCA 181

DB 122 ACGTCAATATGACGTAATGTTCCCATATGACGTAATGAGGACTTTCATGAGCTCA 181

QY 182 TGGGTGAGTATTTACGTAATGCGCGCGGCTGACGCCACGACCCCGCCCATTTG 240

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OM nucleic - nucleic search, using sw model

Run on: December 15, 2002, 05:48:20 ; Search time 68 Seconds

(without alignments)
2503.024 Million cell updates/sec

Title: US-10-059-152-1

Perfect score: 555
Sequence: 1 gagtattataatgaatcaaa.....cggctctgactgaccgctc 555

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 441362 seqs, 153338381 residues

Total number of hits satisfying chosen parameters: 882724

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%
Listing first 45 summaries

Database :

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2: /cgn2_6/ptodata/2/ina/5B.COMB.seq:*
3: /cgn2_6/ptodata/2/ina/6A.COMB.seq:*
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5: /cgn2_6/ptodata/2/ina/PCITS.COMB.seq:*
6: /cgn2_6/ptodata/2/ina/Backfiles1.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	325	58.6	6714	US-09-299-141-6	Sequence 6, Appl1
2	325	58.6	6924	US-09-299-141-9	Sequence 9, Appl1
3	325	58.6	6924	US-09-299-141-10	Sequence 11, Appl1
4	325	58.6	6924	US-09-299-141-11	Sequence 10, Appl1
5	256	46.1	763	US-08-459-493-1	Sequence 1, Appl1
6	256	46.1	930	US-08-029-022-2	Sequence 2, Appl1
7	256	46.1	930	US-08-029-022-4	Sequence 4, Appl1
8	256	46.1	930	US-08-246-376-2	Sequence 2, Appl1
9	256	46.1	930	US-08-246-376-4	Sequence 4, Appl1
10	256	46.1	930	US-07-972-135-2	Sequence 2, Appl1
11	256	46.1	930	US-07-972-135-4	Sequence 4, Appl1
12	256	46.1	930	US-08-256-004-2	Sequence 2, Appl1
13	256	46.1	930	US-09-006-841-2	Sequence 2, Appl1
14	256	46.1	930	US-09-006-841-4	Sequence 4, Appl1
15	256	46.1	930	PCT-US93-05366-2	Sequence 4, Appl1
16	256	46.1	930	PCT-US93-05366-4	Sequence 2, Appl1
17	256	46.1	1078	US-09-310-842-1	Sequence 1, Appl1
18	256	46.1	1318	US-09-310-842-3	Sequence 3, Appl1
19	256	46.1	1417	US-09-310-842-5	Sequence 5, Appl1
20	256	46.1	1645	US-09-310-842-2	Sequence 2, Appl1
21	256	46.1	1870	US-09-310-842-4	Sequence 4, Appl1
22	256	46.1	3987	US-09-082-649B-83	Sequence 83, Appl1
23	256	46.1	3987	US-09-082-649B-84	Sequence 84, Appl1
24	256	46.1	4072	US-09-770-315-4	Sequence 4, Appl1
25	256	46.1	4278	US-09-503-799-2	Sequence 1, Appl1
26	256	46.1	4397	US-09-503-799-1	Sequence 1, Appl1
27	256	46.1	4518	US-09-380-190A-26	Sequence 26, Appl1

28	256	46.1	4627	US-09-054-281-19	Sequence 19, Appl1
29	256	46.1	4665	US-08-948-378A-7	Sequence 7, Appl1
30	256	46.1	4665	US-09-169-425C-7	Sequence 7, Appl1
31	256	46.1	4696	US-08-929-967-15	Sequence 15, Appl1
32	256	46.1	4732	US-07-884-811-1	Sequence 1, Appl1
33	256	46.1	4732	US-07-885-971-1	Sequence 1, Appl1
34	256	46.1	4732	US-08-087-783A-1	Sequence 1, Appl1
35	256	46.1	4732	US-08-194-088B-1	Sequence 1, Appl1
36	256	46.1	4732	US-08-194-088B-1	Sequence 1, Appl1
37	256	46.1	4732	PCT-US93-04648-1	Sequence 1, Appl1
38	256	46.1	4883	US-09-608-730B-21	Sequence 21, Appl1
39	256	46.1	4886	US-09-533-220A-4	Sequence 4, Appl1
40	256	46.1	4915	US-09-173-053-7	Sequence 7, Appl1
41	256	46.1	5158	US-08-929-967-16	Sequence 16, Appl1
42	256	46.1	5261	US-09-770-315-7	Sequence 7, Appl1
43	256	46.1	5452	US-09-130-114-1	Sequence 1, Appl1
44	256	46.1	5552	US-08-155-888-1	Sequence 1, Appl1
45	256	46.1	5585	US-08-305-221-1	Sequence 1, Appl1

ALIGNMENTS

RESULT 1
US-09-299-141-6
Sequence 6, Application US/09299141
Patent No. 6461606
GENERAL INFORMATION:
APPLICANT: FLOTT, TERENCE R.
APPLICANT: SONG, STRONG
APPLICANT: BYRNE, BARRY J.
APPLICANT: MORGAN, MICHAEL
TITLE OF INVENTION: MATERIALS AND METHODS FOR GENE THERAPY
FILE REFERENCE: 4300.011800
CURRENT APPLICATION NUMBER: US/09/299,141
CURRENT FILING DATE: 1999-04-23
EARLIER APPLICATION NUMBER: 60/083,025
EARLIER FILING DATE: 1998-04-24
NUMBER OF SEQ ID NOS: 13
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 6
LENGTH: 6714
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: PLASMID
US-09-299-141-6

Query Match 58.6%; Score 325; DB 4; Length 6714;
Best Local Similarity 74.9%; Pred. No. 8.2e-63;
Matches 543; Conservative 0; Mismatches 10; Indels 172; Gaps 4;

2 AGTATTATAGTATCAATTTACGGGTCATTGCTCATATGCCATATATGAGTCCG 61
|||||
323 AGTATTATAGTATCAATTTACGGGTCATTGCTCATATATGAGTCCG 382
|||||
62 GTTACATTAATCGGTAATTTACGGGTCATTGCTCATATATGAGTCCG 121
|||||
383 GTTACATTAATCGGTAATTTACGGGTCATTGCTCATATATGAGTCCG 442
|||||
122 AGTCAATATAGTATTTACCGATAGTATAGCCATATAGGACTTCCATAGCTCA 181
|||||
443 AGTCAATATAGTATTTACCGATAGTATAGCCATATAGGACTTCCATAGCTCA 502
|||||
182 TGGGTGAGTATTTACCGATAGTATAGCCATATAGGACTTCCATAGCTCA 240
|||||
503 TGGGTGAGTATTTACCGATAGTATAGCCATATAGGACTTCCATAGCTCA 562
|||||
241 AGTACGCCCCCTTACCGATAGTATAGCCATATAGGACTTCCATAGCTCA 276
|||||
563 AGTACGCCCCCTTACCGATAGTATAGCCATATAGGACTTCCATAGCTCA 622


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OY 2 AGTATTATAGTAATCAATTAACGGGGTCATTAGTTCATAGCCCATATATAGAGTTCCG 61
DB 533 AGTTATTAATTAATCAATTAACGGGGTCATTAGTTCATAGCCCATATATAGAGTTCCG 592
OY 62 GTTACATACTTAACGTAATTAAGCCCGGCGGTGACCCAGACAGCCCGCCCATTTG 121
DB 593 GTTACATACTTAACGTAATTAAGCCCGGCGGTGACCCAGACAGCCCGCCCATTTG 652
OY 122 AGCTCAATTAATAGCATATGTTCCCATAGTAACGCCAATAGGACCTTCCATTGACGTCA 181
DB 653 AGCTCAATTAATAGCATATGTTCCCATAGTAACGCCAATAGGACCTTCCATTGACGTCA 712
OY 182 TGGGTGAGTATTAAGGTAACGTCGCA-TTGGCAGTACATCAAGTATATATATGCCA 240
DB 713 TGGGTGAGTATTAAGGTAACGTCGCACTTGGCAGTACATCAAGTATATATATGCCA 772
OY 241 AGTACGCCCCCTATTGACGTCAATGACGTAATAGG----- 276
DB 773 AGTACGCCCCCTATTGACGTCAATGACGTAATAGGCCCCGCGCATTTATGCCAGTAC 832
OY 277 ----- 276
DB 833 ATGACCTTAAGGGAGCTTCTACTTGCGAGTACATCTAGCATATAGTATATATAC 892
OY 277 ----- 276
DB 893 ATGTCGAGGTGAGCCCAAGTTCGTCTACCTCCCATCTCCCGCCCTCCACACC 952
OY 277 -----ATGCAATATTTTGTGACAGCATGGGGC-----GGGGG 310
DB 953 CCAATTTTGTATTTATTTATTTTATTTTATTTTGTGACAGCATATGGGGCGGGGGGGG 1012
OY 311 GGGGGGCGCGCCAGCGGGGGGGGGGGGGGAGAGGGGGGGGGGGGGAGAGGCGAGA 370
DB 1013 GGGGGGCGCGCCAGCGGGGGGGGGGGGGGAGAGGGGGGGGGGGGGAGAGGCGAGA 1072
OY 371 GGTGGGGGGGAGCCCAATCAGAGCGGGGGGGGGGCGGAAAGTTTCTTTATATGGCGAGGGG 430
DB 1073 GGTGGGGGGGAGCCCAATCAGAGCGGGGGGGGGGCGGCGAAAGTTTCTTTATATGGCGAGGGG 1132
OY 431 CGGCGGGGGGGGCGCTATTAATAAGGAGCGCGGGGGGGGGAGTGCCTCG-CCGTG 489
DB 1133 CGGCGGGGGGGGCGCTATTAATAAGGAGCGCGGGGGGGGGAGTGCCTCGAGCGCTG 1192
OY 490 CCTTGGCCCCGTGCGCGCTCGCGCGCTCGCGCGCGCGCGCGCGCGCTCTGACTGAC 549
DB 1193 CCTTGGCCCCGTGCGCGCTCGCGCGCGCGCTCGCGCGCGCGCGCGCGCTCTGACTGAC 1252
OY 550 CGCGT 554
DB 1253 CGCGT 1257

RESULT 4
US-09-299-141-11
: Sequence 11, Application US/09299141
: Patent No. 6461606
: GENERAL INFORMATION:
: APPLICANT: FLOTTE, TERENCE R.
: APPLICANT: SONG, SIONG
: APPLICANT: BYRNE, BARRY J.
: APPLICANT: MORGAN, MICHAEL
: TITLE OF INVENTION: MATERIALS AND METHODS FOR GENE THERAPY
: FILE REFERENCE: 4300.011800
: CURRENT APPLICATION NUMBER: US/09/299,141
: EARLIER FILING DATE: 1999-04-23
: EARLIER APPLICATION NUMBER: 60/083,025
: NUMBER OF SEQ ID NOS: 13
: SOFTWARE: PatentIn Ver. 2.0
: SEQ ID NO 11
: LENGTH: 6924
: TYPE: DNA
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: ORGANISM: Artificial Sequence
: FEATURE:
: OTHER INFORMATION: Description of Artificial Sequence: PLASMID
: OTHER INFORMATION: p43rmeNCB-AF
US-09-299-141-11
Query Match 58.6%; Score 325; DB 4; Length 6924;
Best Local Similarity 74.9%; Pred. No. 8.2e-63;
Matches 543; Conservative 0; Mismatches 10; Indels 172; Gaps 4;

OY 2 AGTATTATAGTAATCAATTAACGGGGTCATTAGTTCATAGCCCATATATAGAGTTCCG 61
DB 533 AGTTATTAATTAATCAATTAACGGGGTCATTAGTTCATAGCCCATATATAGAGTTCCG 592
OY 62 GTTACATACTTAACGTAATTAAGCCCGGCGGTGACCCAGACAGCCCGCCCATTTG 121
DB 593 GTTACATACTTAACGTAATTAAGCCCGGCGGTGACCCAGACAGCCCGCCCATTTG 652
OY 122 AGCTCAATTAATAGCATATGTTCCCATAGTAACGCCAATAGGACCTTCCATTGACGTCA 181
DB 653 AGCTCAATTAATAGCATATGTTCCCATAGTAACGCCAATAGGACCTTCCATTGACGTCA 712
OY 182 TGGGTGAGTATTAAGGTAACGTCGCA-TTGGCAGTACATCAAGTATATATATGCCA 240
DB 713 TGGGTGAGTATTAAGGTAACGTCGCACTTGGCAGTACATCAAGTATATATATGCCA 772
OY 241 AGTACGCCCCCTATTGACGTCAATGACGTAATAGG----- 276
DB 773 AGTACGCCCCCTATTGACGTCAATGACGTAATAGGCCCCGCGCATTTATGCCAGTAC 832
OY 277 ----- 276
DB 833 ATGACCTTAAGGGAGCTTCTACTTGCGAGTACATCTAGCATATAGTATATATAC 892
OY 277 ----- 276
DB 893 ATGTCGAGGTGAGCCCAAGTTCGTCTACCTCCCATCTCCCGCCCTCCACACC 952
OY 277 -----ATGCAATATTTTGTGACAGCATGGGGC-----GGGGG 310
DB 953 CCAATTTTGTATTTATTTATTTTATTTTATTTTGTGACAGCATATGGGGCGGGGGGGG 1012
OY 311 GGGGGGCGCGCCAGCGGGGGGGGGGGGGGAGAGGGGGGGGGGGGGAGAGGCGAGA 370
DB 1013 GGGGGGCGCGCCAGCGGGGGGGGGGGGGGAGAGGGGGGGGGGGGGAGAGGCGAGA 1072
OY 371 GGTGGGGGGGAGCCCAATCAGAGCGGGGGGGGGGCGGAAAGTTTCTTTATATGGCGAGGGG 430
DB 1073 GGTGGGGGGGAGCCCAATCAGAGCGGGGGGGGGGCGGCGAAAGTTTCTTTATATGGCGAGGGG 1132
OY 431 CGGCGGGGGGGGCGCTATTAATAAGGAGCGCGGGGGGGGGAGTGCCTCG-CCGTG 489
DB 1133 CGGCGGGGGGGGCGCTATTAATAAGGAGCGCGGGGGGGGGAGTGCCTCGAGCGCTG 1192
OY 490 CCTTGGCCCCGTGCGCGCTCGCGCGCTCGCGCGCGCGCGCGCGCTCTGACTGAC 549
DB 1193 CCTTGGCCCCGTGCGCGCTCGCGCGCGCGCTCGCGCGCGCGCGCGCGCTCTGACTGAC 1252
OY 550 CGCGT 554
DB 1253 CGCGT 1257

RESULT 5
US-08-459-493-1
: Sequence 1, Application US/08459493
: Patent No. 6030638
: GENERAL INFORMATION:
: APPLICANT: Brigham, Kenneth
: APPLICANT: Conarty, Jon T.
: APPLICANT: Canonico, Angelo
: APPLICANT: Meyrick, Barbara
: TITLE OF INVENTION: PLASMID FOR IN VIVO EXPRESSION
```

;; TITLE OF INVENTION: PROSTAGLANDIN SYNTHASE
;; NUMBER OF SEQUENCES: 3
;; CORRESPONDENCE ADDRESS:
;; ADDRESSEE: Tilton, Fallon, Lungmus & Chestnut
;; STREET: 100 S. Wacker Drive - Suite 960
;; CITY: Chicago
;; STATE: Illinois
;; COUNTRY: U.S.A.
;; ZIP: 60606-4002
;; COMPUTER READABLE FORM:
;; MEDIUM TYPE: Floppy disk
;; OPERATING SYSTEM: IBM PC compatible
;; SOFTWARE: Patentin Release #1.0, Version #1.25
;; CURRENT APPLICATION DATA:
;; APPLICATION NUMBER: US/08/459,493
;; FILING DATE:
;; CLASSIFICATION: 435
;; PRIOR APPLICATION DATA:
;; APPLICATION NUMBER: US 08/080,221
;; FILING DATE: 21-JUNE-1993
;; APPLICATION NUMBER: US 07/746,941
;; FILING DATE: 19-AUG-1991
;; ATTORNEY/AGENT INFORMATION:
;; NAME: Kohn, Kenneth I.
;; REGISTRATION NUMBER: 30,955
;; TELECOMMUNICATION INFORMATION:
;; TELEPHONE: (313) 456-8000
;; INFORMATION FOR SEQ ID NO: 1:
;; SEQUENCE CHARACTERISTICS:
;; LENGTH: 763 base pairs
;; TYPE: nucleic acid
;; STRANDEDNESS: double
;; TOPOLOGY: circular
;; MOLECULE TYPE: DNA (genomic)
;; HYPOTHETICAL: NO
;; ANTI-SENSE: NO
;; FEATURE:
;; NAME/KEY: Promoter
;; LOCATION: 1..763
;; US-08-459-493-1

Query Match 46.1%; Score 256; DB 3; Length 763;
Best Local Similarity 97.8%; Pred. No. 7.7e-48;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

0Y 2 AGTTATTAATAGTAAATTCAGGAGGATTCATAGCCCAATATGAGTTCCGC 61
|||
DB 180 AGTTATTAATAGTAAATTCAGGAGGATTCATAGCCCAATATGAGTTCCGC 239
|||
0Y 62 GTTACATACTTACGTAATTTGCGCGGCTGACCGCCCAACGACCCCGCCCATG 121
|||
DB 240 GTTACATACTTACGTAATTTGCGCGGCTGACCGCCCAACGACCCCGCCCATG 299
|||
0Y 122 AGCTCAATATAGCTATGTTCCCATAGTAAAGGCAATGAGGACTTTCATTGAGCTCA 181
|||
DB 300 AGCTCAATATAGCTATGTTCCCATAGTAAAGGCAATGAGGACTTTCATTGAGCTCA 359
|||
0Y 182 TGGGTGAGATTTTACGTAACCTGCCA-TTGGCAGTACATCAAGTATCATATGCA 240
|||
DB 360 TGGGTGAGATTTTACGTAACCTGCCA-TTGGCAGTACATCAAGTATCATATGCA 419
|||
0Y 241 AGTACGCCCTATTGAGCTCAATGACGTAATGG 276
|||
DB 420 AGTACGCCCTATTGAGCTCAATGACGTAATGG 455
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RESULT 6
US-08-029-022-2
; Sequence 2, Application US/08029022
; Patent No. 5641662
; GENERAL INFORMATION:

;; APPLICANT: Debs, Robert J.
;; APPLICANT: Zhu, Ming
;; TITLE OF INVENTION: TRANSFECTION OF LUNG VIA AEROSOLIZED
;; TITLE OF INVENTION: TRANSFECTED DELIVERY
;; NUMBER OF SEQUENCES: 4
;; CORRESPONDENCE ADDRESS:
;; ADDRESSEE: Fish & Richardson
;; STREET: 2200 Sand Hill Road, Suite 100
;; CITY: Menlo Park
;; STATE: California
;; COUNTRY: U.S.A.
;; ZIP: 94025
;; COMPUTER READABLE FORM:
;; MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
;; COMPUTER: IBM PS/2 Model 502 or 55SX
;; OPERATING SYSTEM: MS-DOS (Version 5.0)
;; SOFTWARE: WordPerfect (Version 5.1)
;; CURRENT APPLICATION DATA:
;; APPLICATION NUMBER: US/08/029,022
;; FILING DATE: 19930310
;; CLASSIFICATION: 514
;; PRIOR APPLICATION DATA:
;; APPLICATION NUMBER: 809,291
;; FILING DATE: 17 DEC 91
;; APPLICATION NUMBER: 972,135
;; FILING DATE: 05 NOV 92
;; APPLICATION NUMBER: PCT/US92/11008
;; FILING DATE: 17 DEC 92
;; ATTORNEY/AGENT INFORMATION:
;; NAME: Rae-Venter, Barbara
;; REGISTRATION NUMBER: 32,750
;; REFERENCE/DOCKET NUMBER: 05935/008001
;; TELECOMMUNICATION INFORMATION:
;; TELEPHONE: (415) 854-5277
;; TELEFAX: (415) 854-0875
;; INFORMATION FOR SEQ ID NO: 2:
;; SEQUENCE CHARACTERISTICS:
;; LENGTH: 930
;; TYPE: nucleic acid
;; STRANDEDNESS: single
;; TOPOLOGY: linear
;; US-08-029-022-2

Query Match 46.1%; Score 256; DB 1; Length 930;
Best Local Similarity 97.8%; Pred. No. 7.9e-48;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

0Y 2 AGTTATTAATAGTAAATTCAGGAGGATTCATAGCCCAATATGAGTTCCGC 61
|||
DB 157 AGTTATTAATAGTAAATTCAGGAGGATTCATAGCCCAATATGAGTTCCGC 216
|||
0Y 62 GTTACATACTTACGTAATTTGCGCGGCTGACCGCCCAACGACCCCGCCCATG 121
|||
DB 217 GTTACATACTTACGTAATTTGCGCGGCTGACCGCCCAACGACCCCGCCCATG 276
|||
0Y 122 AGCTCAATATAGCTATGTTCCCATAGTAAAGGCAATGAGGACTTTCATTGAGCTCA 181
|||
DB 277 AGCTCAATATAGCTATGTTCCCATAGTAAAGGCAATGAGGACTTTCATTGAGCTCA 336
|||
0Y 182 TGGGTGAGATTTTACGTAACCTGCCA-TTGGCAGTACATCAAGTATCATATGCA 240
|||
DB 337 TGGGTGAGATTTTACGTAACCTGCCA-TTGGCAGTACATCAAGTATCATATGCA 396
|||
0Y 241 AGTACGCCCTATTGAGCTCAATGACGTAATGG 276
|||
DB 397 AGTACGCCCTATTGAGCTCAATGACGTAATGG 432
|||

RESULT 7
US-08-029-022-4
; Sequence 4, Application US/08029022
; Patent No. 5641662

GENERAL INFORMATION:

APPLICANT: Debs, Robert J.

APPLICANT: Zhu, Ning

TITLE OF INVENTION: TRANSECTION OF LUNG VIA AEROSOLIZED

TITLE OF INVENTION: TRANSENE DELIVERY

NUMBER OF SEQUENCES: 4

CORRESPONDENCE ADDRESS:

ADDRESSEE: Fish & Richardson

STREET: 2200 Sand Hill Road, Suite 100

CITY: Menlo Park

STATE: California

COUNTRY: U.S.A.

ZIP: 94025

COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5" Diskette, 1.44 MB

COMPUTER: IBM PS/2 Model 502 or 55SX

OPERATING SYSTEM: MS-DOS (Version 5.0)

SOFTWARE: WordPerfect (Version 5.1)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/029,022

FILING DATE: 19930310

CLASSIFICATION: 514

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 809,291

FILING DATE: 17 DEC 91

APPLICATION NUMBER: 972,135

FILING DATE: 05 NOV 92

APPLICATION NUMBER: PCT/US92/11008

FILING DATE: 17 DEC 92

ATTORNEY/AGENT INFORMATION:

NAME: Rae-Ventler, Barbara

REGISTRATION NUMBER: 32,750

REFERENCE/DOCKET NUMBER: 05935/008001

TELECOMMUNICATION INFORMATION:

TELEPHONE: (415) 854-5277

TELEFAX: (415) 854-0875

TELEX: 200154

INFORMATION FOR SEQ ID NO: 4:

SEQUENCE CHARACTERISTICS:

LENGTH: 930

TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

US-08-029-022-4

Query Match 46.1%; Score 256; DB 1; Length 930;

Best Local Similarity 97.8%; Pred. No. 7.9e-48;

Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 2 AGTTATTAATAGTATCAATTAACGGGGGTCATTAGTTCATAGCCCATATATGAGTTCCGC 61
DB 157 AGTTATTAATAGTATCAATTAACGGGGGTCATTAGTTCATAGCCCATATATGAGTTCCGC 216
QY 62 GTTACATACTAGCGTAATATGCGCCCGGCTGACCGCCCAAGAGACCCCGCCCATG 121
DB 217 GTTACATACTAGCGTAATATGCGCCCGGCTGACCGCCCAAGAGACCCCGCCCATG 276
QY 122 ACGTCAATATAGCGTAATATGCGCCCGGCTGACCGCCCAAGAGACCCCGCCCATG 181
DB 277 ACGTCAATATAGCGTAATATGCGCCCGGCTGACCGCCCAAGAGACCCCGCCCATG 336
QY 182 TGGGTGAGTATTTACGTAACCTGCCA-TTGGCAGTACATCAAGTATATATAGCA 240
DB 337 TGGGTGAGTATTTACGTAACCTGCCA-TTGGCAGTACATCAAGTATATATAGCA 396
QY 241 ACTAGCCCCCTATTGAGCTCATGAGGTAATGG 276
DB 397 ACTAGCCCCCTATTGAGCTCATGAGGTAATGG 432

RESULT 8
US-08-246-376-2
Sequence 2, Application US/08246376

Patent No. 5827703

GENERAL INFORMATION:

APPLICANT: Robert J. Debs

APPLICANT: Ning Zhu

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR IN VIVO

TITLE OF INVENTION: GENE THERAPY

NUMBER OF SEQUENCES: 4

CORRESPONDENCE ADDRESS:

ADDRESSEE: Fish & Richardson

STREET: 2200 Sand Hill Road, Suite 100

CITY: Menlo Park

STATE: California

COUNTRY: U.S.A.

COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5" Diskette, 1.44 MB

COMPUTER: IBM PS/2 Model 502 or 55SX

OPERATING SYSTEM: MS-DOS (Version 5.0)

SOFTWARE: WordPerfect (Version 5.1)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/246,376

FILING DATE:

CLASSIFICATION: 800

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 07/992,687

FILING DATE: December 17, 1992

APPLICATION NUMBER: 07/927,200

FILING DATE: August 6, 1992

APPLICATION NUMBER: 07/894,498

FILING DATE: June 4, 1992

ATTORNEY/AGENT INFORMATION:

NAME: Barbara Rae-Ventler

REGISTRATION NUMBER: 32,750

REFERENCE/DOCKET NUMBER: 05935/005US1

TELECOMMUNICATION INFORMATION:

TELEPHONE: (415) 854-5277

TELEFAX: (415) 854-0875

TELEX: 200154

INFORMATION FOR SEQ ID NO: 2:

SEQUENCE CHARACTERISTICS:

LENGTH: 930

TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

US-08-246-376-2

Query Match 46.1%; Score 256; DB 1; Length 930;

Best Local Similarity 97.8%; Pred. No. 7.9e-48;

Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 2 AGTTATTAATAGTATCAATTAACGGGGGTCATTAGTTCATAGCCCATATATGAGTTCCGC 61
DB 157 AGTTATTAATAGTATCAATTAACGGGGGTCATTAGTTCATAGCCCATATATGAGTTCCGC 216
QY 62 GTTACATACTAGCGTAATATGCGCCCGGCTGACCGCCCAAGAGACCCCGCCCATG 121
DB 217 GTTACATACTAGCGTAATATGCGCCCGGCTGACCGCCCAAGAGACCCCGCCCATG 276
QY 122 ACGTCAATATAGCGTAATATGCGCCCGGCTGACCGCCCAAGAGACCCCGCCCATG 181
DB 277 ACGTCAATATAGCGTAATATGCGCCCGGCTGACCGCCCAAGAGACCCCGCCCATG 336
QY 182 TGGGTGAGTATTTACGTAACCTGCCA-TTGGCAGTACATCAAGTATATATAGCA 240
DB 337 TGGGTGAGTATTTACGTAACCTGCCA-TTGGCAGTACATCAAGTATATATAGCA 396
QY 241 ACTAGCCCCCTATTGAGCTCATGAGGTAATGG 276
DB 397 ACTAGCCCCCTATTGAGCTCATGAGGTAATGG 432

RESULT 9
US-08-246-376-4

Sequence 4, Application US/08246376
Patent No. 5827703
GENERAL INFORMATION:
APPLICANT: Robert J. Debs
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR IN VIVO
TITLE OF INVENTION: GENE THERAPY
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 2200 Sand Hill Road, Suite 100
CITY: Menlo Park
STATE: California
COUNTRY: U.S.A.
ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: IBM PS/2 Model 502 or 55SX
OPERATING SYSTEM: MS-DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/246,376
FILING DATE:
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/992,687
FILING DATE: December 17, 1992
APPLICATION NUMBER: 07/927,200
FILING DATE: August 6, 1992
APPLICATION NUMBER: 07/894,498
FILING DATE: June 4, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Barbara Rae-Venter
REGISTRATION NUMBER: 32,750
REFERENCE/DOCKET NUMBER: 05935/005051
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 854-5277
TELEFAX: (415) 854-0875
TELEX: 200154
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 930
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
US-08-246-376-4

Query Match 46.1% Score 256; DB 1; Length 930;
Best Local Similarity 97.8%; Pred. No. 7.9e-48;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;
DB 2 AGTATTAAATAGTAAATACGAGGTCATTATGTCATAGCCCATATATGAGTTCCGC 61
DB 157 AGTATTAAATAGTAAATACGAGGTCATTATGTCATAGCCCATATATGAGTTCCGC 216
QY 62 GTTACATTAATAGTAAATAGGCGCGCGGTGACGCCCAAGACCCCGCCCATTTG 121
DB 217 GTTACATTAATAGTAAATAGGCGCGCGGTGACGCCCAAGACCCCGCCCATTTG 276
QY 122 AGTCAATTAATAGTAAATAGGCGCGCGGTGACGCCCAAGACCCCGCCCATTTG 181
DB 277 AGTCAATTAATAGTAAATAGGCGCGCGGTGACGCCCAAGACCCCGCCCATTTG 336
QY 182 TGGGTGAGTAAATAGTAAATAGGCGCGCGGTGACGCCCAAGACCCCGCCCATTTG 240
DB 337 TGGGTGAGTAAATAGTAAATAGGCGCGCGGTGACGCCCAAGACCCCGCCCATTTG 396
QY 241 AGTACGCCCTATTGACGTCAATGACGTAATAGG 276
DB 397 AGTACGCCCTATTGACGTCAATGACGTAATAGG 432

RESULT 10

US-07-972-135-2
Sequence 2, Application US/07972135
Patent No. 5858784
GENERAL INFORMATION:
APPLICANT: Robert J. Debs
TITLE OF INVENTION: EXPRESSION OF CLONED
TITLE OF INVENTION: GENES IN THE
TITLE OF INVENTION: LUNG BY AEROSOL - AND
TITLE OF INVENTION: LIPOSOE-BASED
TITLE OF INVENTION: DELIVERY
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 2200 Sand Hill Road,
CITY: Menlo Park
STATE: California
COUNTRY: U.S.A.
ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: IBM PS/2 Model 502 or
COMPUTER: 55SX
OPERATING SYSTEM: MS-DOS (Version 5.0)
SOFTWARE: WordPerfect (Version
SOFTWARE: 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/972,135
FILING DATE: No. 5858784ember 5, 1992
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/809,291
FILING DATE: December 17, 1991
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Barbara Rae-Venter
REGISTRATION NUMBER: 32,750
REFERENCE/DOCKET NUMBER: 05935/007051
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 854-5277
TELEFAX: (415) 854-0875
TELEX: 200154
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 930
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
US-07-972-135-2

Query Match 46.1% Score 256; DB 2; Length 930;
Best Local Similarity 97.8%; Pred. No. 7.9e-48;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;
DB 2 AGTATTAAATAGTAAATACGAGGTCATTATGTCATAGCCCATATATGAGTTCCGC 61
DB 157 AGTATTAAATAGTAAATACGAGGTCATTATGTCATAGCCCATATATGAGTTCCGC 216
QY 62 GTTACATTAATAGTAAATAGGCGCGCGGTGACGCCCAAGACCCCGCCCATTTG 121
DB 217 GTTACATTAATAGTAAATAGGCGCGCGGTGACGCCCAAGACCCCGCCCATTTG 276
QY 122 AGTCAATTAATAGTAAATAGGCGCGCGGTGACGCCCAAGACCCCGCCCATTTG 181
DB 277 AGTCAATTAATAGTAAATAGGCGCGCGGTGACGCCCAAGACCCCGCCCATTTG 336
QY 182 TGGGTGAGTAAATAGTAAATAGGCGCGCGGTGACGCCCAAGACCCCGCCCATTTG 240
DB 337 TGGGTGAGTAAATAGTAAATAGGCGCGCGGTGACGCCCAAGACCCCGCCCATTTG 396
QY 241 AGTACGCCCTATTGACGTCAATGACGTAATAGG 276

Db 397 AGTACGCCCCCTATTGACGTCAATGACGTAATGG 432

RESULT 11
US-07-972-135-4

GENERAL INFORMATION:
APPLICANT: Robert J. Debs
TITLE OF INVENTION: EXPRESSION OF CLONED
GENES IN THE
TITLE OF INVENTION: LONG BT AEROSOL - AND
TITLE OF INVENTION: LIPOSOE-BASED
TITLE OF INVENTION: DELIVERY
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:

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? COMPUTER READABLE FORM:
? MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
? COMPUTER: IBM PS/2 Model 50Z or
? OPERATING SYSTEM: MS-DOS (Version 5.0)
? SOFTWARE: Wordperfect (Version
? E 1

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Query Match	46.1%	Score 256	DB 2	Length 930
Best Local Similarity	97.8%	Pred. No.	7.9e-48	
Matches 270	Conservative	0	Mismatches 5	Indels 1
				Gaps 1

Db 337 TGGGTGAGTATTACGGTAACCTGCCACCTGGCAGTACATCAAGTGTATCATATGCCA 396

```

QY      241 AGTACGCCCCCTATTGACGTCAATGACGGTAAATGG 276
          |||||
Db      397 AGTACGCCCCCTATTGACGTCAATGACGGTAAATGG 432

```

RESULT 12
US-08-256-004-2
; Sequence 2, Application US/08256004

GENERAL INFORMATION:
APPLICANT: Robert J. Debs
APPLICANT: Ning Zhu
TITLE OF INVENTION: IN VIVO TRANSFECTION WITH A CTR CODING
TITLE OF INVENTION: SEQUENCE
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooley Godward Castro Huddleson & Tatum
STREET: 5 Palo Alto Square
CITY: Palo Alto
STATE: California

```

1  COMPUTER READABLE FORM:
2  MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
3  COMPUTER: IBM PC compatible
4  OPERATING SYSTEM: PC-DOS/MS-DOS
5  SOFTWARE: Patent In Release #1.0, Version #1.25
6  CURRENT APPLICATION DATA:
7  APPLICATION NUMBER: US/08/256,004
8  FILING DATE: August 22, 1994

```

US-08-256-004-2

Query Match	256;	Score	256;	DB 3;	length	930;
Best Local Similarity	97.88;	Pred. No.	7.9e-48;			
Matches 270;	Conservative	0;	Mismatches	5;	Indels	1;
					Gaps	1;

DB 277 ACCTCAATATATGACGTATGTCCCATAGTACGCCCATAGGACTTCCATTACGTCGA 336
|||
QY 182 TGGGTGAGTATTATAGGTAACCTGCCA-TTGGCAGTACATCAAGTATCATATGCCA 240
|||
DB 337 TGGGTGAGTATTATAGGTAACCTGCCA-TTGGCAGTACATCAAGTATCATATGCCA 396
|||
QY 241 AGTACGCCCCCTATGACGTCAATGACGGTAATGG 276
|||
DB 397 AGTACGCCCCCTATGACGTCAATGACGGTAATGG 432
|||

RESULT 13

US-09-006-841-2
; Sequence 2, Application US/09006841
; Patent No. 6468798
; GENERAL INFORMATION:
; APPLICANT: Robert J. Debs
; TITLE OF INVENTION: EXPRESSION OF CLONED
; TITLE OF INVENTION: GENES IN THE
; TITLE OF INVENTION: LONG BY AEROSOL - AND
; TITLE OF INVENTION: LIPOSOME-BASED
; TITLE OF INVENTION: DELIVERY
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: Fish & Richardson
; STREET: 2200 Sand Hill Road,
; CITY: Menlo Park
; STATE: California
; COUNTRY: U.S.A.
; ZIP: 94025
; COMPUTER READABLE FORM:
; MEDIUM TYPE: 3.5" Diskette, 1.44 MB
; COMPUTER: IBM PS/2 Model 502 or 55SX
; OPERATING SYSTEM: MS-DOS (Version 5.0)
; SOFTWARE: WordPerfect (Version 5.1)
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/006,841
; FILING DATE:
; CLASSIFICATION:
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: US/07/972,135
; FILING DATE: No. 6468798, December 5, 1992
; APPLICATION NUMBER: 07/809,291
; FILING DATE: December 17, 1991
; ATTORNEY/AGENT INFORMATION:
; NAME: Barbara Rae-Venter
; REGISTRATION NUMBER: 32,750
; REFERENCE/DOCKET NUMBER: 05935/007US1
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 854-5277
; TELEFAX: (415) 854-0875
; TELEX: 200154
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 930
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; US-09-006-841-2

Query Match 46.1%; Score 256; DB 4; Length 930;
Best Local Similarity 97.8%; Pred. No. 7.9e-48;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 2 AGTTATTAATAGTAATCAATTTACGGGGTCAATTGTTTCATAGCCCATATATGAGTTCCG 61
|||
DB 157 AGTTATTAATAGTAATCAATTTACGGGGTCAATTGTTTCATAGCCCATATATGAGTTCCG 216
|||
QY 62 GTTACATTAATCGGTAATATGGCCGCGCTGACGCCCAAGCAGACCCCGCCATTG 121
|||
DB 217 GTTACATTAATCGGTAATATGGCCGCGCTGACGCCCAAGCAGACCCCGCCATTG 276
|||

QY 122 ACCTCAATATATGACGTATGTCCCATAGTACGCCCATAGGACTTCCATTACGTCGA 181
|||
DB 277 ACCTCAATATATGACGTATGTCCCATAGTACGCCCATAGGACTTCCATTACGTCGA 336
|||
QY 182 TGGGTGAGTATTATAGGTAACCTGCCA-TTGGCAGTACATCAAGTATCATATGCCA 240
|||
DB 337 TGGGTGAGTATTATAGGTAACCTGCCA-TTGGCAGTACATCAAGTATCATATGCCA 396
|||
QY 241 AGTACGCCCCCTATGACGTCAATGACGGTAATGG 276
|||
DB 397 AGTACGCCCCCTATGACGTCAATGACGGTAATGG 432
|||

RESULT 14

US-09-006-841-4
; Sequence 4, Application US/09006841
; Patent No. 6468798
; GENERAL INFORMATION:
; APPLICANT: Robert J. Debs
; TITLE OF INVENTION: EXPRESSION OF CLONED
; TITLE OF INVENTION: GENES IN THE
; TITLE OF INVENTION: LONG BY AEROSOL - AND
; TITLE OF INVENTION: LIPOSOME-BASED
; TITLE OF INVENTION: DELIVERY
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: Fish & Richardson
; STREET: 2200 Sand Hill Road,
; CITY: Menlo Park
; STATE: California
; COUNTRY: U.S.A.
; ZIP: 94025
; COMPUTER READABLE FORM:
; MEDIUM TYPE: 3.5" Diskette, 1.44 MB
; COMPUTER: IBM PS/2 Model 502 or 55SX
; OPERATING SYSTEM: MS-DOS (Version 5.0)
; SOFTWARE: WordPerfect (Version 5.1)
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/006,841
; FILING DATE:
; CLASSIFICATION:
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: US/07/972,135
; FILING DATE: No. 6468798, December 5, 1992
; APPLICATION NUMBER: 07/809,291
; FILING DATE: December 17, 1991
; ATTORNEY/AGENT INFORMATION:
; NAME: Barbara Rae-Venter
; REGISTRATION NUMBER: 32,750
; REFERENCE/DOCKET NUMBER: 05935/007US1
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 854-5277
; TELEFAX: (415) 854-0875
; TELEX: 200154
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 930
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; US-09-006-841-4

Query Match 46.1%; Score 256; DB 4; Length 930;
Best Local Similarity 97.8%; Pred. No. 7.9e-48;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 2 AGTTATTAATAGTAATCAATTTACGGGGTCAATTGTTTCATAGCCCATATATGAGTTCCG 61
|||
DB 157 AGTTATTAATAGTAATCAATTTACGGGGTCAATTGTTTCATAGCCCATATATGAGTTCCG 216
|||
QY 62 GTTACATTAATCGGTAATATGGCCGCGCTGACGCCCAAGCAGACCCCGCCATTG 121
|||

GenCore version 5.1.3
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OM nucleic - nucleic search, using sw model

Run on: December 15, 2002, 05:46:45 ; Search time 2426 Seconds

(Without alignments)
3705.068 Million cell updates/sec

Title: US-10-059-152-1

Perfect score: 555

Sequence: 1 gagtataatagtaacaa.....cggctctgactgacgcgctc 555

Scoring table:

IDENTITY_NUC
Gapop 10.0 , Gapept 1.0

Searched: 16154066 seqs, 8097743376 residues

Total number of hits satisfying chosen parameters: 32308132

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

EST:*
1: em_estba:*
2: em_esthum:*
3: em_estlin:*
4: em_estlin:*
5: em_estlin:*
6: em_estlin:*
7: em_estlin:*
8: em_estlin:*
9: gb_est1:*
10: gb_est2:*
11: gb_est3:*
12: gb_est4:*
13: gb_est5:*
14: gb_est6:*
15: em_estfun:*
16: em_estfun:*
17: gb_gss:*
18: em_gss_hum:*
19: em_gss_inv:*
20: em_gss_pln:*
21: em_gss_vrt:*
22: em_gss_fun:*
23: em_gss_mam:*
24: em_gss_mus:*
25: em_gss_other:*
26: em_gss_pro:*
27: em_gss_rtd:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	254.4	45.8	569	12	BC063359 H3006C07-3
2	254.4	45.8	578	9	AA09312 EST03748
3	231.6	41.7	675	10	AV645339 AV645339
4	224.4	40.4	642	10	AV645353 AV645353
5	215.4	38.8	672	10	AV681528 AV681528
6	213.4	38.5	645	10	AV681458 AV681458

Result No.	Score	Query Match	Length	ID	Description
7	210	37.8	783	10	AV681502
8	207.8	37.4	676	10	AV681477
9	205.8	37.1	619	10	AV645377
10	199.2	35.9	525	14	BO247946
11	195.4	35.2	401	12	BC899290
12	195.2	35.2	821	10	AV681519
13	194	35.0	806	10	AV681464
14	192.2	34.6	807	10	AV681462
15	191.6	34.5	672	10	AV681510
16	183.8	33.1	674	10	AV681487
17	183	33.0	674	10	AV681514
18	181.8	32.8	803	10	AV681483
19	181.4	32.7	619	10	AV645379
20	181.4	32.7	811	10	AV681506
21	181	32.6	790	10	AV681497
22	180.8	32.6	620	10	AV645369
23	179.6	32.4	816	10	AV681523
24	178.8	32.2	683	10	AV681491
25	169	30.5	641	10	AV645404
26	164.6	29.7	630	10	AV645321
27	159	28.6	269	14	BO757691
28	156.6	28.2	802	10	AV681486
29	155.8	28.1	180	13	BI823701
30	153	27.6	633	10	AV681527
31	153	27.6	649	14	BO248366
32	152	27.4	648	14	BO247997
33	152	27.4	648	14	BO248418
34	152	27.4	648	14	BO248913
35	151	27.2	647	14	BO248311
36	151	27.2	647	14	BO248458
37	151	27.2	649	14	BO247933
38	150	27.0	646	14	BO247762
39	150	27.0	646	14	BO248341
40	149.8	27.0	686	10	AV681472
41	149	26.8	645	14	BO247960
42	148.2	26.7	619	10	AV645389
43	148	26.7	644	14	BO247963
44	146	26.3	642	14	BO247949
45	145	26.1	641	14	BO248159

ALIGNMENTS

RESULT 1
LOCUS BC063359
DEFINITION H3006C07-3 NIA Mouse 15K CDNA Clone Set Mus musculus EST 26-JAN-2001
ACCESSION BC063359
VERSION BC063359.1 GI:12545922
KEYWORDS
SOURCE
ORGANISM Mus musculus
REFERENCE
AUTHORS Kargul, G.J., Dudekula, D.B., Qian, Y., Lim, M.K., Jaradat, S.A., Tanaka, T.S., Carter, M.G. and Ko, M.S.H.
TITLE Verification and initial annotation of NIA mouse 15K CDNA clone set
JOURNAL Unpublished (2001)
COMMENT Contact: George J. Kargul
Laboratory of Genetics
National Institute on Aging/National Institutes of Health
333 Cassell Drive, Suite 4000, Baltimore, MD 21224-6820, USA
Email: cdna@nigms.nih.gov
This clone set has been freely distributed to the community. Please visit <http://lgsun.grc.nia.nih.gov/cDNA/15k.html> for details.
Plate: H306 Row: C Column: 07
Seq primer: -21M13 Forward
High quality sequence stop: 569
POLY-A-No.

FEATURES

Location/Qualifiers

source 1.569

/organism="Mus musculus"

/strain="C57BL/6J"

/db_xref="nlsest:H3006C07-3"

/db_xref="taxon:10090"

/clone="H3006C07"

/clone_1lb="NIA Mouse 15K cDNA Clone Set"

/sex="Clones arrayed from a variety of cDNA libraries"

/dev_stage="Clones arrayed from a variety of cDNA libraries"

/lab_host="DH10B"

/note="Vector: pSPORT1; Site.1: SalI; Site.2: NotI; This clone is among a rearranged set of 15,247 clones from 11 embryo cDNA libraries (including preimplantation stage embryos from unfertilized egg to blastocyst, embryonic part of E7.5 embryos, extraembryonic part of E7.5 embryos and E12.5 female mesonephros/gonad) and one newborn ovary cDNA library. Average insert size 1.5 kb. All source libraries are cloned unidirectionally with Oligo(dT) -Not primers. References include: (1) Genome-wide expression profiling of mid-gestation placenta and embryo using a 15,000 mouse developmental cDNA microarray, 2000, Proc. Natl. Acad. Sci. U S A. 97: 9127-9132; (2) Large-scale cDNA analysis reveals phased gene expression patterns during preimplantation mouse development, 2000; Development, 127: 1737-1749; (3) Genome-wide mapping of unselected transcripts from extraembryonic tissue of 7.5-day mouse embryos reveals enrichment in the t-complex and under-representation on the X chromosome, 1998, Hum Mol Genet 7: 1967-1978."

BASE COUNT 150 a 136 c 130 g 153 t

ORIGIN

Query Match 45.8%; Score 254.4; DB 12; Length 569;

Best Local Similarity 97.5%; Pred. No. 3.5e-45;

Matches 269; Conservative 0; Mismatches 6; Indels 1; Gaps 1;

OY 2 AGTTATTAAATAGTAATCAATTAAGGGGTCAATAGTTCATAGCCCATATATGAGTTCGCC 61

DB 222 AGTATTAAATAGTAATCAATTAAGGGGTCAATAGTTCATAGCCCATATATGAGTTCGCC 281

OY 62 GTTACATTAATAGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 121

DB 282 GTTACATTAATAGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 341

OY 122 ACCTGATTAATAGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 181

DB 342 ACCTGATTAATAGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 401

OY 182 TGGGTGAGATTAATAGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 240

DB 402 TGGGTGAGATTAATAGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 461

OY 241 AGTACGGCCCTATTTGACGTCAATGACGTAATATGG 276

DB 462 AGTACGGCCCTATTTGACGTCAATGACGTAATATGG 497

RESULT 2

AA409312 578 bp mRNA linear EST 26-AUG-1998

LOCUS EST03748 Mouse 7.5 dpc embryo ectoplacental cone cDNA library Mus

DEFINITION musculus cDNA clone C0039E08 3' similar to cloning vector pCI-neo,

AA409312 mammalian expression vector,, score = 2136, mRNA sequence.

ACCESSION AA409312

VERSION AA409312.1 GI:2068663

KEYWORDS EST.

SOURCE house mouse.

ORGANISM Mus musculus

REFERENCE 1 (bases 1 to 578)

Authors Ko,M.S.H., Threlk,T.A., Horton,J.H., Wang,X, Cui,Y., Wang,X., Pryor

JOURNAL Unpublished (1997)

COMMENT Contact: Ko MSH

Center for Molecular Medicine and Genetics

Wayne State University

5047 Guillen Mall, Detroit, MI 48202

Tel: 3135776708

Fax: 3135776200

Email: mskocmb.biosci.wayne.edu

Cloning vector pCI-neo, mammalian expression vector,, score = 2136

Seq primer: M13 Forward.

Location/Qualifiers

source 1.578

/organism="Mus musculus"

/strain="C57BL/6J"

/db_xref="ATCC (Inhost):1364275"

/db_xref="taxon:10090"

/clone="C0039E08"

/clone_1lb="Mouse 7.5 dpc embryo ectoplacental cone cDNA library"

/sex="unknown"

/tissue_type="ectoplacental cone"

/dev_stage="embryonic day 7.5 postconception"

/lab_host="DH10B"

/note="Organ: embryo; Vector: pSPORT1 (Life Technologies); Site.1: SalI; Site.2: NotI; Total RNAs were extracted from ectoplacental cone of 7.5-dpc embryos. The double-stranded cDNA was synthesized from total RNAs with an Oligo(dT) primer. The library was constructed by Minoru S. H. Ko."

BASE COUNT 153 a 138 c 130 g 154 t 3 others

ORIGIN

Query Match 45.8%; Score 254.4; DB 9; Length 578;

Best Local Similarity 97.5%; Pred. No. 3.5e-45;

Matches 269; Conservative 0; Mismatches 6; Indels 1; Gaps 1;

OY 2 AGTTATTAAATAGTAATCAATTAAGGGGTCAATAGTTCATAGCCCATATATGAGTTCGCC 61

DB 234 AGTATTAAATAGTAATCAATTAAGGGGTCAATAGTTCATATAGCCCATATATGAGTTCGCC 293

OY 62 GTTACATTAATAGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 121

DB 294 GTTACATTAATAGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 353

OY 122 ACCTGATTAATAGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 181

DB 354 ACCTGATTAATAGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 413

OY 182 TGGGTGAGATTAATAGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 240

DB 414 TGGGTGAGATTAATAGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 473

OY 241 AGTACGGCCCTATTTGACGTCAATGACGTAATATGG 276

DB 474 AGTACGGCCCTATTTGACGTCAATGACGTAATATGG 509

RESULT 3

AV645339 675 bp mRNA linear EST 15-JAN-2002

LOCUS AV645339/c

DEFINITION AV645339 GLA Homo sapiens cDNA clone GLAAC04 3', mRNA sequence.

ACCESSION AV645339

VERSION AV645339.1 GI:9866353

KEYWORDS EST.

SOURCE human.

ORGANISM Homo sapiens

REFERENCE 1 (bases 1 to 675)

Authors Nakashima,H.,

Systematic analyses of mouse genes expressed in embryo implantation

site

AUTHORS	Xu, X., Huang, J., Xu, Z., Qian, B., Zhu, Z., Yan, Q., Cai, T., Zhang, X., Xiao, H., Qu, J., Liu, F., Huang, Q., Cheng, Z., Li, N., Du, J., Hu, W., Shen, K., Lu, G., Fu, G., Zhong, M., Xu, S., Gu, W., Huang, W., Zhao, X., Hu, G., Gu, J., Chen, Z., and Han, Z.
TITLE	Insight into hepatocellular carcinogenesis at transcriptome level by comparing gene expression profiles of hepatocellular carcinoma with those of corresponding noncancerous liver
JOURNAL	Proc. Natl. Acad. Sci. U.S.A. 98 (26), 15089-15094 (2001)
COMMENT	21625106 Contact: Zeguang Han Chinese National Human Genome Center at Shanghai 351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, Pudong, Shanghai 201203, P. R. China Tel: 86-21-50801919(ex. 45) Fax: 86-21-50801922 Email: hanzg@hgc.sh.cn This clone is available at CHGC in Shanghai.
FEATURES	Location/Qualifiers 1..675 /organism="Homo sapiens" /db_xref="taxon:9606" /clone="GLAAC04" /clone_lib="GLA" /tissue_type="corresponding non cancerous liver tissue" /dev_stage="Adult" /lab_host="SODR" /note="Vector: pbluescript sk(-); Site_1: EcoRI; Site_2: XhoI"
BASE COUNT	159 a 166 c 177 g 169 t 4 others
ORIGIN	
Query Match	41.7%; Score 231.6; DB 10; Length 675;
Best Local Similarity	93.0%; Pred. No. 2.9e-40;
Matches 251; Conservative	0; Mismatches 18; Indels 1; Gaps 1;
OY	8 TAATGTAATCAATTACGGGGTCAATTAGTTCATACCCATATATGAGTCCGCGTTACA 67 Db 673 TTATGGATCAANATACGGGGTCAATTAGTTCATATATGAGTCCGCGTTACA 614 68 TAACCTAGCGTAAATGCGCGCGGCTGACGCCCAACGACCCCGCCATTGAGGTCA 127 Db 613 TACTTACGGGTAATAGCGCCCGCTGCTGACCGCCCAACGACCCCGCCATTGAGGTCA 554 128 ATATAGAGGTATGTCCCATAGTAACGCCAATAGGAGACTTCCATTGACGTCAATGGGTG 187 553 ATATAGAGGTATGTCCCATAGTAACGCCAATAGGAGACTTCCATTGAGGTCAATGGGTG 494 OY 188 GAGTATTTACGGTAAACGTGCCA-TTGGCAGTACATCAAGTGTATCATATGCCAAGTAGC 246 Db 493 GAGTATTTACGGTAAACGTGCCA-TTGGCAGTACATCAAGTGTATCATATGCCAAGTAGC 434 OY 247 CCCCTATTTAGCGTCAATGATGCGTAAATGG 276 Db 433 CCCCTATTTAGCGTCAATGATGCGTAAATGG 404
RESULT 4	
LOCUS	AV645353 642 bp mRNA linear EST 15-JAN-2002
DEFINITION	AV645353 GLA Homo sapiens cDNA clone GLAAD06 3', mRNA sequence.
ACCESSION	AV645353
VERSION	AV645353.1 GI:9866367
KEYWORDS	EST.
SOURCE	human.
ORGANISM	Homo sapiens
REFERENCE	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominiidae; Homo.
AUTHORS	1 (bases 1 to 642) Xiao, H., Huang, J., Xu, Z., Qian, B., Zhu, Z., Yan, Q., Cai, T., Zhang, X., Xiao, H., Qu, J., Liu, F., Huang, Q., Cheng, Z., Li, N., Du, J., Hu, W., Shen, K., Lu, G., Fu, G., Zhong, M., Xu, S., Gu, W., Huang, W., Zhao, X., Hu, G., Gu, J., Chen, Z., and Han, Z.
TITLE	Insight into hepatocellular carcinogenesis at transcriptome level

JOURNAL MEDLINE COMMENT	by comparing gene expression profiles of hepatocellular carcinoma with those of corresponding noncancerous liver Proc. Natl. Acad. Sci. U.S.A. 98 (26), 15089-15094 (2001)
21625106	Contact: Zeguang Han
	Chinese National Human Genome Center at Shanghai
	351 Guo Shoujing Road, Zhanjiang Hi-Tech Park, Pudong, Shanghai
	201203, P. R. China
	Tel: 86-21-50801919 (ex.45)
	Fax: 86-21-50801922
	Email: hanzgzhc@sh.cn
	This clone is available at CHGC in Shanghai.
FEATURES	Location/Qualifiers
source	1. .642
	/organism="Homo sapiens"
	/db_xref="taxon:9606"
	/clone="GUAAD06"
	/clone_lib="GTA"
	/issue_type="corresponding non cancerous liver tissue"
	/dev_stage="Adult"
	/db_host="SOLR"
	/note="Vector: pbluescript sk(-); Site_1: EcoRI; Site_2: XhoI"
BASE COUNT	150 a 160 c 168 g 160 t 4 others
ORIGIN	
Query Match	40.4%; Score 224.4; DB 10; Length 642;
Best Local Similarity	93.7%; Pred. No. 1e-38;
Matches 253; Conservative	0; Mismatches 15; Indels 2; Gaps 2;
QY 8	TAATAGTAAATCAATTAACGGGGTCATTAGTTCATAGCCCATATATAGAGATCCGGTTACA 67
Db 635	TTATAGGATCAATATACGGGGTCANTAGTTCATAGCCCATATATAGAGATCCGGTTACA 576
QY 68	TAACTTACGGTAAATTTGGCCCGCGGCTGACGCCCAAGACCCCGCCCAATTGACGTCA 127
Db 575	T-ACCTTACGGTAAAGGGCGCCCTGGCTGGACGCCCAAGACCCCGCCCAATTGACGTCA 517
QY 128	ATAATGACGTATGTCCCATAGTAAAGCCCATATAGGAGATTTCCATATGACGTCAATGGGTG 187
Db 516	ATAATGACGTATGTTCCTATAGTAAAGCCCATATAGGAGATTTCCATATGACGTCAATGGGTG 457
QY 188	GAGTATTTACGGTAAACTGCCA-TTGGCAGTACATCAAGTATATATATCCCAAGTACG 246
Db 456	AAGATATTAGCGTAAACGCCACCTGGCACTATACATCAATCAAGTATATATCCCAAGTACG 397
QY 247	CCCCCTATTGACGTCAATGACGTAAATGG 276
Db 396	CCCCCTATTGACGTCAATGACGTAAATGG 367
RESULT 5	
AV681528/c	672 bp mRNA linear EST 16-JAN-2002
LOCUS	AV681528
DEFINITION	AV681528 GK A Homo sapiens cDNA clone GRAAB04 5', mRNA sequence.
ACCESSION	AV681528
VERSION	AV681528.1 GI:10283391
KEYWORDS	EST.
SOURCE	human.
ORGANISM	Homo sapiens
REFERENCE	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
AUTHORS	Mammalia; Eutheria; Primates; Catarrhini; Homiidae; Homo.
	1 (bases 1 to 672)
	Xu, X., Huang, J., Xu, Z., Qian, B., Zhu, Z., Yan, Q., Cai, T., Zhang, X.,
	Xiao, H., Qu, J., Liu, F., Huang, Q., Cheng, Z., Li, N., Du, J., Hu, W.,
	Shen, K., Lu, G., Fu, G., Zhong, M., Xu, S., Gu, W., Huang, W., Zhao, X.,
	Hu, G., Gu, J., Chen, Z., and Han, Z.
	Insight into hepatocellular carcinogenesis at transcriptome level
	by comparing gene expression profiles of hepatocellular carcinoma
	with those of corresponding noncancerous liver
	Proc. Natl. Acad. Sci. U.S.A. 98 (26), 15089-15094 (2001)
JOURNAL	21625106
MEDLINE	Contact: Zeguang Han
COMMENT	Chinese National Human Genome Center at Shanghai


```

/db_xref="taxon:9606"
/clone="GKAAH01"
/clone_lib="GKA"
/tissue_type="hepatocellular carcinoma"
/dev_stage="Adult"
/lab_host="SOLR"
/Note="Vector: pBluescript sk(-); Site_1: EcoRI; Site_2:
XhoI"
BASE COUNT      198 a      191 c      191 g      201 t      2 others
ORIGIN
Query Match      37.8%; Score 210; DB 10; Length 783;
Best Local Similarity 89.5%; Pred. No. 1.3e-35;
Matches 247; Conservative 0; Mismatches 27; Indels 2; Gaps 2;

OY 2 AGTATTAAATGATCAATTAAGGGGCTATTGTCATATGACCCATATATGAGTCCGC 61
    |||||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||
DB 720 AGTATTAAATGATCAATTAAGGGGCTC-TTGCTTCAAAAGCCATTAATGAAGTCCGC 662
    |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||
OY 62 GTTACATTAAGTAAATTTGGCCGCGCTGACCCGCAAGACCCGCCCATTTG 121
    |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||
DB 661 GTTACATTAAGTAAATTTGGCCGCGCTGACCCGCTGACCCGCTGACCCGCCCATTTG 602
    |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||
OY 122 AGCTCAATTAATGACGATGTTCCCATATGATACGCCCAATGAGACTTTCATGACGTCAA 181
    |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||
DB 601 ACCTCAATTAATGACGATGTTCCCATATGATACGCCCAATGAGACTTTCATGACGTCAA 542
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OY 182 TGGGTGAGATTTTAACTGCTAACTGCCCA-TTGGCAGTACATCAAGTATTCATATGCCA 240
    |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||
DB 541 TGGGTGAGATTTTAACTGCTAACTGCCCATTTGGCAGTACATCAAGTATTCATATGCCA 482
    |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||
OY 241 AGTACGCCCTTATGACGCTAATGACGGTAAATGG 276
    |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||
DB 481 AGTACGCCCTTATGACGCTAATGACGGTAAATGG 446
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RESULT 8
AV681477/c      676 bp      mRNA      linear      EST 16-JAN-2002
LOCUS      AV681477 GKA Homo sapiens cDNA clone GKAAAB12 5', mRNA sequence.
DEFINITION      AV681477
ACCESSION      AV681477
VERSION      AV681477.1 GI:10283340
KEYWORDS      EST.
SOURCE      human.
ORGANISM      Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.
REFERENCE      1 (bases 1 to 676)
AUTHORS      Xu,X., Huang,J., Xu,Z., Qian,B., Zhu,Z., Yan,Q., Cai,T., Zhang,X.,
Xiao,H., Qu,J., Liu,F., Huang,Q., Cheng,Z., Li,N., Du,J., Hu,W.,
Shen,K., Lu,G., Fu,G., Zhong,M., Xu,S., Gu,W., Huang,W., Zhao,X.,
Hu,G., Gu,Y., Chen,Z. and Han,Z.
TITLE      Insight into hepatocellular carcinogenesis at transcriptome level
by comparing gene expression profiles of hepatocellular carcinoma
with those of corresponding noncancerous liver
JOURNAL      Proc. Natl. Acad. Sci. U.S.A. 98 (26), 15089-15094 (2001)
MEDLINE      21625106
COMMENT      Contact: Zeguang Han
Chinese National Human Genome Center at Shanghai
351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, Pudong, Shanghai
201203, P. R. China
Tel: 86-21-50801919(ex.45)
Fax: 86-21-50801922
Email: hanzg@hgc.sh.cn
This clone is available at CHGC in Shanghai.
FEATURES
source
1. 676
/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="GKAAAB12"
/clone_lib="GKA"
/tissue_type="hepatocellular carcinoma"
/dev_stage="Adult"

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/lab_host="SOLR"
/Note="Vector: pBluescript sk(-); Site_1: EcoRI; Site_2:
XhoI"
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Best Local Similarity 91.6%; Pred. No. 4e-35;
Matches 240; Conservative 0; Mismatches 20; Indels 2; Gaps 2;

OY 16 ATCAATTACGGGGTCAATTAGTTCATAGCCCATATATGAGTCCGCTTACATTAATAC 75
    |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||
DB 672 ATCAATTACGGGGTCAATTAGTTCATAGCCCATATATGAGTCCGCTTACATTAATAC 613
    |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||
OY 76 GGTAAATTTGGCCGCGCTGACCCGCAAGACCCGCCCATTTGACGTCAATTAATGAC 135
    |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||
DB 612 GGTAAATTTGGCCGCGCTGACCCGCAAGACCCGCCCATTTGACGTCAATTAATGAC 553
    |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||
OY 136 GTATGTCCCATTAATGACCCCAATGAGACTTTCATTAATGACGTCAATGAGTGAATTT 195
    |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||
DB 552 GTATGTCCCATTAATGACCCCAATGAGACTTTCATTAATGACGTCAATGAGTGAATTT 494
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    |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||  |||||||
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RESULT 9
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LOCUS      AV645377 GLA Homo sapiens cDNA clone GLAABF06 3', mRNA sequence.
DEFINITION      AV645377
ACCESSION      AV645377
VERSION      AV645377.1 GI:9866391
KEYWORDS      EST.
SOURCE      human.
ORGANISM      Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.
REFERENCE      1 (bases 1 to 619)
AUTHORS      Xu,X., Huang,J., Xu,Z., Qian,B., Zhu,Z., Yan,Q., Cai,T., Zhang,X.,
Xiao,H., Qu,J., Liu,F., Huang,Q., Cheng,Z., Li,N., Du,J., Hu,W.,
Shen,K., Lu,G., Fu,G., Zhong,M., Xu,S., Gu,W., Huang,W., Zhao,X.,
Hu,G., Gu,Y., Chen,Z. and Han,Z.
TITLE      Insight into hepatocellular carcinogenesis at transcriptome level
by comparing gene expression profiles of hepatocellular carcinoma
with those of corresponding noncancerous liver
JOURNAL      Proc. Natl. Acad. Sci. U.S.A. 98 (26), 15089-15094 (2001)
MEDLINE      21625106
COMMENT      Contact: Zeguang Han
Chinese National Human Genome Center at Shanghai
351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, Pudong, Shanghai
201203, P. R. China
Tel: 86-21-50801919(ex.45)
Fax: 86-21-50801922
Email: hanzg@hgc.sh.cn
This clone is available at CHGC in Shanghai.
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[illegible]

ACCESSION AV681510
 VERSION AV681510.1 GI:10283373
 KEYWORDS EST.
 SOURCE human.
 ORGANISM Homo sapiens
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.
 REFERENCE 1 (bases 1 to 672)
 AUTHORS Xu,X., Huang,J., Xu,Z., Qian,B., Zhu,Z., Yan,Q., Cai,T., Zhang,X., Xiao,H., Qu,J., Liu,F., Huang,Q., Cheng,Z., Li,N., Du,J., Hu,W., Shen,K., Lu,G., Zhong,H., Xu,S., Gu,W., Huang,W., Zhao,X., Hu,G., Gu,J., Chen,Z., and Han,Z.
 TITLE Insignt into hepatocellular carcinogenesis at transcriptome level by comparing gene expression profiles of hepatocellular carcinoma with those of corresponding noncancerous liver
 JOURNAL Proc. Natl. Acad. Sci. U.S.A. 98 (26), 15089-15094 (2001)
 MEDLINE 21625106
 COMMENT Contact: Zequang Han
 Chinese National Human Genome Center at Shanghai
 351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, Pudong, Shanghai 201203, P. R. China
 Tel: 86-21-50801919(ex.45)
 Fax: 86-21-50801922
 Email: hanzq@chgc.sh.cn
 This clone is available at CHGC in Shanghai.
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 /note="Vector: pbluescript sk(-); Site_1: EcoRI; Site_2: XhoI"
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 Best Local Similarity 85.6%; Pred. No. 1.2e-31;
 Matches 220; Conservative 0; Mismatches 36; Indels 1; Gaps 1.
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 Oy 81 ATTGGCCGCCGCGGTGACGCCCAACGACCCCGCCCATTTGACGTCAATATGAGAGTAT 140
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 Oy 141 TTCCCATAGTAAAGCCCATATGAGGACTTTTCATTTGACGTCAATATGAGGATTTTACG 200
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 Db 492 AAAGTGGCCA-TTGGCAGTACATCAAGTGTATCATATGCGCAAGTACGCCCTTATGAG 433
 Oy 260 TCAATGACGTAATATG 276
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 Db 432 TCAATGACGTAATATG 416

GenCore version 5.1.3
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OM nucleic - nucleic search, using sw model

Run on: December 15, 2002, 05:41:10 ; Search time 3127 Seconds
(without alignments) 5165.352 Million cell updates/sec

Title: US-10-059-152-1

Perfect score: 555
Sequence: 1 gatttataatgtaataca.....cggctctgactgaccgcgyc 555

Scoring table: IDENTITY_NDC
Gapop 10.0 , Gapept 1.0

Searched: 2054640 seqs, 14551402878 residues

Total number of hits satisfying chosen parameters: 4109280

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :

- 1: gb_ba:*
- 2: gb_htg:*
- 3: gb_in:*
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- 22: em_ov:*
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- 27: em_sts:*
- 28: em_un:*
- 29: em_vl:*
- 30: em_htg_hum:*
- 31: em_htg_in:*
- 32: em_htg_other:*
- 33: em_htg_mu:*
- 34: em_htg_pl:*
- 35: em_htg_rod:*
- 36: em_htg_syn:*
- 37: em_htg_vrt:*
- 38: em_sy:*
- 39: em_htgo_hum:*
- 40: em_htgo_mu:*
- 41: em_htgo_other:*

Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

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6	309.4	55.7	6148	6 AX468470	AX468470 Sequence
7	257	46.3	599	6 E51990	E51990 Promoter, r
8	256	46.1	590	6 E59416	E59416 Signal pept
9	256	46.1	777	6 BD000145	BD000145 Vector ha
10	256	46.1	777	6 I05430	I05430 Sequence 16
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17	256	46.1	930	6 AR050544	AR050544 Sequence
18	256	46.1	930	6 AR050546	AR050546 Sequence
19	256	46.1	930	6 AR094363	AR094363 Sequence
20	256	46.1	930	6 I49834	I49834 Sequence 2
21	256	46.1	930	6 I49836	I49836 Sequence 4
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23	256	46.1	1078	6 A92081	A92081 Sequence 5
24	256	46.1	1318	6 A92080	A92080 Sequence 4
25	256	46.1	1417	6 A92077	A92077 Sequence 1
26	256	46.1	1645	6 A92078	A92078 Sequence 2
27	256	46.1	1767	6 AX402407	AX402407 Sequence
28	256	46.1	1848	14 HEHCMP1	X03922 Human clyom
29	256	46.1	1870	6 A92079	A92079 Sequence 3
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34	256	46.1	3602	12 AF327894	AF327894 Cloning v
35	256	46.1	3610	6 AX030966	AX030966 Sequence
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37	256	46.1	3903	6 AX076478	AX076478 Sequence
38	256	46.1	3987	6 AR182910	AR182910 Sequence
39	256	46.1	3987	6 AR182911	AR182911 Sequence
40	256	46.1	4050	12 AF053407	AF053407 Expressio
41	256	46.1	4072	6 AX205074	AX205074 Sequence
42	256	46.1	4187	12 AF151087	AF151087 Cloning v
43	256	46.1	4201	6 AX076477	AX076477 Sequence
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ALIGNMENTS

RESULT 1	LOCUS	DEFINITION	ACCESSION	VERSION	KEYWORDS	SOURCE	ORGANISM	REFERENCE	AUTHORS	TITLE	JOURNAL
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COMMENT OS Artificial Sequence
 PN JP 2001000188-A/3
 PD 09-JAN-2001
 PF 22-JUN-1999 JP 1999174804
 PR MAYUMI KUBOMURA, SHUJI SAITO
 PI C12N15/09, A61K31/00, A61K39/02, A61K39/17, A61K39/215,
 PC C07K14/125
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 BASE COUNT 109 a 156 c 191 g 105 t
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 Matches 553; Conservative 0; Mismatches 0; Indels 3; Gaps 2;

QY 2 AGTTATTAATAGTAATCAATTAAGGGGTCATTGTTCCATAGACCCCATATATAGAGTCCGC 61
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 DB 61 GTTACATACTTAAGGTAATTTGGCCCGGCTGACCCGCCAAGACCCCGCCCATTTG 119
 QY 122 AGCTCAATTAATGACGATGTGTCATAGTACGCCCAATAGGAGCTTCCATAGAGTCGAA 181
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 QY 182 TGGGTGAGTAATTAAGGTAATTTGGCCCATTTGGCAGTACATCAAGTATCATATGCCAA 241
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 DEFINITION AF334827
 ACCESSION AF334827.1 GI:12965137
 VERSION
 KEYWORDS Cloning vector pTurbo-Cre.
 SOURCE

ORGANISM Cloning vector pTurbo-Cre
 artificial sequences; vectors.
 REFERENCE 1 (bases 1 to 5894)
 AUTHORS Lu,Z.H., Graubert,T.A. and Ley,T.J.
 TITLE Direct Submission
 JOURNAL Submitted (10-JAN-2001) Division of Oncology, Section of Stem Cell
 Biology, Washington University School of Medicine, 660 South Euclid
 Avenue, Campus Box 8007, St. Louis, MO 63110-1093, USA
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 QY 122 AGCTCAATTAATGACGATGTGTCATAGTACGCCCAATAGGAGCTTCCATAGAGTCGAA 181
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 QY 182 TGGGTGAGTAATTAAGGTAATTTGGCCCATTTGGCAGTACATCAAGTATCATATGCCAA 240
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				Gaps 5
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DB	198	GTTACATAACTATAGGTAATTTGGCCCCCGCTGACGCCCAAGSACCCCGCCATTG	257	
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DB	258	ACGTCAATATAGCATGTATGCCATAGTAAAGCCCAATAGGAGCTTTCATTGACGTCA	317	
OY	182	TGGGAGAGTATTTACGGTAAACTGCCA-TTGGAGATACATCAAGTATCATATGCA	240	
DB	318	TGGGAGAGTATTTACGGTAAACTGCCA-TTGGAGATACATCAAGTATCATATGCA	377	
OY	241	AGTAGCCCCCTATGAGCTCAATGACGGTAATGG-----	276	
DB	378	AGTAGCCCCCTATGAGCTCAATGACGGTAATGG-----	437	
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DB	618	GGGGGCGCGCGCAGAGCGGGGGCGGGGCGAGGGGGGGGGCGAGGCGGAGAG	677	
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DB	678	GTCGGCGCGCAGCATCAGAGCGGGGCTCCGAAAGTTTCTTTTATGGCGAGGCGGC	737	
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DB	738	GGCGGGGGGGGCGCTATATAAAGCGAAGGGGCGGGGGGGGAGGCTGCGCTGGC	795	
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DB	855	CGT 857		
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SV	BD005270.1			
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DT	08-FEB-2002 (Rel. 70, Last updated, Version 1)			
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XX	JP 03074399-T/2.			

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OS		artificial sequence.	
CC			
XX			
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RA	Ishino F., Miyoshi N., Ishino T., Yokoyama M., Wakana S.;		
RT	"Mammalian model for diabetes";		
RL	Patent number JP03074399-T/2, 12-JAN-2001.		
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CC	PJ 18-AUG-2000 JP 2000005546		
CC	PR 20-OCT-1999 JP 99P 298273		
CC	PI FUJIHOSHII ISHINO,NAOKI MIYOSHI,TOMOKO ISHINO,MINESUKE KOROZAMA,		
CC	PI SHIGEHARU WAKANA		
CC	PC A01K07/027,C12N15/12,C12Q1/68,C12Q1/02,A6IK45/00,A6IP3/10,		
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	Matches 539; Conservative 0; Mismatches 11; Indels 173; Gaps		
QY	2 AGTTAATTAAGTAAATCAATTACGGGTCATTACTTAGTACCACCATATATGAGGTGCCG 61		
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QY	62 GTTACATACTTACCGTAAATTTGGCCCCGCCTCACC GCCCACGAGCCCCCCCATTG 121		
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ACCESSION	AX468471			
VERSION	AX468471.1	GI:21901307		
KEYWORDS				
SOURCE				
ORGANISM				
REFERENCE	1			
AUTHORS	Siebel, C. and Brennan, T.J.			
TITLE	Methods of producing cells and animals comprising targeted gene modifications			
JOURNAL	Patent: WO 0222834-A 14 21-MAR-2002; Deltagen, Inc. (US)			
FEATURES				
source		Location/Qualifiers		
		1..5100		
		/organism="synthetic construct"		
		/db_xref="taxon:32630"		
		/note="Construct Sequence"		
BASE COUNT	1124 a	1357 c	1475 g	1144 t
ORIGIN				
Query Match		55.74;	Score 309.4;	DB 6; Length 5100;
Best Local Similarity		74.6%;	Pred. No. 1.8e-51;	
Matches 539; Conservative		0; Mismatches 11;	Indels 173; Gaps	5;
QY	2	AGTTTATATAGTATATCATTCACGAGGGGTCATATGTTCAATGAGCCATATATGAGTTCCGC	61	
Db	59	AGTTTATATAGTATATCATTCACGAGGGGTCATATGTTCAATGAGCCATATATGAGTTCCGC	118	
QY	62	GTTAATACACTTACGGTAATTTGGCCCGCGGCTGACCGCCACGACCCCGCCCATTTG	121	
Db	119	GTTAATACACTTACGGTAATTTGGCCCGCGGCTGACCGCCACGACCCCGCCCATTTG	178	
QY	122	ACGCAATAAAGACGATGTTCCCATATGTAAGCAATGAGCAATGAGCAATGAGCAATGAGCAAT	181	
Db	179	ACGCAATAAAGACGATGTTCCCATATGTAAGCAATGAGCAATGAGCAATGAGCAATGAGCAAT	238	
QY	182	TGGGAGGAGTATTTACGGTAAACTGCGCA--TTGGCAGTACATCAAGTATCATATGCGCA	240	
Db	239	TGGGAGGAGTATTTACGGTAAACTGCGCACTTGGCAGTACATCAAGTATCATATGCGCA	298	
QY	241	AGTACGCCCCCTATTGACGCTCAATACGGTAATGG-----	276	
Db	299	AGTACGCCCCCTATTGACGCTCAATACGGTAATGG-----	358	
QY	277	-----	276	
Db	359	ATGACCTTACGCGGACTTCTCTACTGCGAGTACATCTACGTATAGTACATGCGCTATTACC	418	
QY	277	-----	276	

Db	2755	AGTAGGCCCCCATTTAGACGTGAATGACGGTAAATGCGCCGCGCTGGCATTTAGCCAGTAGC	2814
Oy	277	-----	276
Db	2815	ATGACCTTAGGGAGCTTCTCCTTGGCAGACATCTAAGTATTAGTCATCTATTATACC	2874
Oy	277	-----	276
Db	2875	ATGGTTCGAGGTGAGGCCCCAGCGTTCTGCTTACTCTCCCCATCTCCCCCTCCCCACG	2934
Oy	277	-----ATGACGATATTTTGTGCAGCGATGGGGGCG--GGGGGGG	311
Db	2935	CCCAATTTTGTATTTATTTATTTTATTTTAAATTTTGTGACGAGATGGGGGCGGGGGGGG	2994
Oy	312	GGGGGGCGCGCCAGCGGGGGCGGGGGCGGAGAGGGCGGGGGCGGAGGCGGAGAG	371
Db	2995	GGGGGGCGCGCCAGCGGGGGCGGGGGCGGAGAGGGCGGGGGCGGAGGCGGAGAG	3054
Oy	372	GGGGGGCGCGCCAGCGGGGGCGGGGGCGGAGAGGGTCTTTTAAAGGAGGGCGG	431
Db	3055	GGGGGGCGCGCCAGCGGGGGCGGGGGCGGAGAGGGTCTTTTAAAGGAGGGCGG	3114
Oy	432	GGGGGGCGCGCCATTTAAAGCAAGCGCGGGGGCGGGGGAGTCTCGCGCTGCG	491
Db	3115	GGGGGGCGCGCCATTTAAAGCAAGCGCGGGGGCGGGGGAGTCTCT--GCTTGGC	3172
Oy	492	TTTGGCCCGCGTCCCGCTCCGCGCGCGCGCGCGCGCGCGCGCGCTCTGACTGACG	551
Db	3173	TTTGGCCCGCGTCCCGCTCCGCGCGCGCGCGCGCGCGCGCGCGCTCTGACTGACG	3231
Oy	552	CGT 554	
Db	3232	CGT 3234	
RESULT 7			
LOCUS	E51990	599 bp	DNA
DEFINITION	Promoter, recombinant containing the same and utilization thereof.		linear PAT 13-AUG-2002
ACCESSION	E51990		
VERSION	E51990.1 GI:18629551		
KEYWORDS	JP 2001000188-A/5.		
SOURCE	Human herpesvirus 5.		
ORGANISM	Human herpesvirus 5.		
REFERENCE	1. Kubomura, M. and Saito, S.		
AUTHORS	Promoter, recombinant containing the same and utilization thereof		
TITLE	Patent: JP 2001000188-A 5 09-JAN-2001;		
JOURNAL	NIPPON ZEON CORP		
COMMENT	OS Cytomegalovirus		
	PN JP 2001000188-A/5		
	PD 09-JAN-2001		
	PF 22-JUN-1999 JP 1999174804		
	PR MAYUMI KUBOMURA, SHUJI SAITO		
	PI C12N15/09, A61K31/00, A61K39/02, A61K39/17, A61K39/215,		
	PC C07K14/125,		
	PC C07K14/165, C07K14/30, C12N7/00, C12N15/00		
FEATURES			
Source	Key	Location/Qualifiers	
	FT	1. 599	
	FT	Location/Qualifiers	
	1. 599	/organism='Cytomegalovirus'	
BASE COUNT	152 a	144 c	144 g 159 t
ORIGIN			
Query Match	46.3%	Score 257;	DB 6; Length 599;
Best Local Similarity	97.8%	Pred No. 4.5e-41;	

Matches 271; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 1 GAGTATTATATAGTCAATTAACGGGGTCATTAGTTCATAGCCCATATATATGAGTTCCG 60
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 DB 6 GAGTATTATATAGTCAATTAACGGGGTCATTAGTTCATAGCCCATATATATGAGTTCCG 65
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QY 61 CGTTACATTAACCTAGCAATTAATGCGCGCTGACCGCCACGACCCCGCCCAT 120
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 DB 66 CATTACATTAACCTAGCAATTAATGCGCGCTGACCGCCACGACCCCGCCCAT 125
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QY 121 GAGTCATTAATGACGATGTTCCCATAGTAAGCCCAATAGGAGCTTCCATTGACGTCA 180
 |||||||
 DB 126 GAGTCATTAATGACGATGTTCCCATAGTAAGCCCAATAGGAGCTTCCATTGACGTCA 185
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QY 181 ATGGGTGAGATTAATGAGTAACTGCCCCA-TTGGCAGTACATCAATGATCATATATGCC 239
 |||||||
 DB 186 ATGGGTGAGATTAATGAGTAACTGCCCCA-TTGGCAGTACATCAATGATCATATATGCC 245
 |||||||

QY 240 AAGTACGCCCCCTATGAGTCAATGAGCGTAATATG 276
 |||||||
 DB 246 AAGTACGCCCCCTATGAGTCAATGAGCGTAATATG 282
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RESULT 8
 E59416
 LOCUS E59416 590 bp DNA linear PAT 31-JAN-2002
 DEFINITION Signal peptide.
 ACCESSION E59416
 VERSION E59416.1 GI:18622549
 KEYWORDS JP 2000354490-A/3.
 SOURCE unidentified.
 ORGANISM unidentified.
 REFERENCE 1 (bases 1 to 590)
 AUTHORS Hawkins, R. and Nakamura, M.
 TITLE Signal peptide
 JOURNAL Patent: JP 2000354490-A 3 26-DEC-2000;
 TOTOTA MOTOR CORP
 OS Cauliflower mosaic virus promoter
 PN JP 2000354490-A/3
 PD 26-DEC-2000
 PE 15-JUN-1999 JP 1999168271
 PR RICHARD HAWKINS, MICHIO NAKAMURA
 PC C12N15/09, C07K14/61, C12N1/13, C12P21/02, C12N15/00 CC
 FH key Location/Qualifiers
 FT source 1..590
 FEATURES Location/Qualifiers
 source 1..590
 /organism="unidentified"
 /db_xref="taxon:32644"

BASE COUNT 149 a 143 c 141 g 157 t

ORIGIN

Query Match 46.1%; Score 256; DB 6; Length 590;
 Best Local Similarity 97.8%; Pred. No. 7.1e-41;
 Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

DB 182 TGGGTGAGATTAATGAGTAACTGCCACTTGACGATACATCAAGTATCATATGCCA 241
 |||||||

QY 241 AGTACGCCCCCTATTGACGTCATATGACGTAATATG 276
 |||||||

DB 242 AGTACGCCCCCTATTGACGTCATATGACGTAATATG 277
 |||||||

RESULT 9
 BD000145
 LOCUS BD000145 777 bp DNA linear PAT 31-JAN-2002
 DEFINITION Vector having stabilized sequence and eucaryotic host cell.
 ACCESSION BD000145
 VERSION BD000145.1 GI:18623224
 KEYWORDS JP 2000308497-A/3.
 SOURCE unidentified.
 ORGANISM unidentified.
 REFERENCE 1 (bases 1 to 777)
 AUTHORS Goman, C.M.
 TITLE Vector having stabilized sequence and eucaryotic host cell
 JOURNAL Patent: JP 2000308497-A 3 07-NOV-2000;
 GENETIC INC
 OS Unidentified
 PN JP 2000308497-A/3
 PD 07-NOV-2000
 PE 17-APR-2000 JP 2000115248
 PR 12-SEP-1986 US 907195, 09-JUL-1987 US 071674 P1
 CORNELIUS MAXIN GOMAN
 PC C12N15/09, C12N15/10, C12N15/00, C12N5/00
 CC
 FH key Location/Qualifiers
 FT source 1..773
 FEATURES Location/Qualifiers
 source 1..777
 /organism="unidentified"
 /db_xref="taxon:32644"

BASE COUNT 188 a 205 c 190 g 194 t

ORIGIN

Query Match 46.1%; Score 256; DB 6; Length 777;
 Best Local Similarity 97.8%; Pred. No. 7e-41;
 Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 2 AGTATTAAATAGTAATCAATTAACGGGGTCATTAGTTCATAGCCCATATATGAGTTCCG 61
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DB 31 AGTATTAAATAGTAATCAATTAACGGGGTCATTAGTTCATAGCCCATATATGAGTTCCG 90
 |||||||

QY 62 GTTACATTAATTAACGTAATATGCGCGCGCTGACCGCCCAACGACCCCGCCCAT 121
 |||||||

DB 91 GTTACATTAATTAACGTAATATGCGCGCGCTGACCGCCCAACGACCCCGCCCAT 150
 |||||||

QY 122 ACATCAATTAATGAGTATGTTCCCATAGTAAGCCCAATAGGAGCTTCCATTGACGTCA 181
 |||||||

DB 151 ACATCAATTAATGAGTATGTTCCCATAGTAAGCCCAATAGGAGCTTCCATTGACGTCA 210
 |||||||

QY 182 TGGGTGAGATTAATGAGTAACTGCCA-TTGGCAGTACATCAAGTATCATATGCCA 240
 |||||||

DB 211 TGGGTGAGATTAATGAGTAACTGCCA-TTGGCAGTACATCAAGTATCATATGCCA 270
 |||||||

QY 241 AGTACGCCCCCTATTGACGTCATATGACGTAATATG 276
 |||||||

DB 271 AGTACGCCCCCTATTGACGTCATATGACGTAATATG 306
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RESULT 10
 I05430
 LOCUS I05430 777 bp DNA linear PAT 02-DEC-1994
 DEFINITION Sequence 16 from Patent EP 0260148.
 ACCESSION I05430
 VERSION I05430.1 GI:591075
 KEYWORDS
 SOURCE Unknown.

ORGANISM Unknown.
REFERENCE 1 (bases 1 to 777)
AUTHORS Gorman,C.M.
TITLE Improved recombinant expression method, vector and transformed cells
JOURNAL Patent: EP 0260148-A2 16 16-MAR-1988;
FEATURES location/Qualifiers
source 1..777
BASE COUNT 188 a 205 c 190 g 194 t
ORIGIN

Query Match 46.1%; Score 256; DB 6; Length 777;
Best Local Similarity 97.8%; Pred. No. 7e-41;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

Db 2 AGTATTATATAGTATCAATTAACGGGGTCATTAGTTCATATAGAGTTCCGC 61
Db 31 AGTATTATATAGTATCAATTAACGGGGTCATTAGTTCATATATAGAGTTCCGC 90
Qy 62 GTTACATTAACCTTACGTAATTTGGCCCGCGCTGACGCCCAAGAGCCCGCCCATTTG 121
Db 91 GTTACATTAACCTTACGTAATTTGGCCCGCGCTGACGCCCAAGAGCCCGCCCATTTG 150
Qy 122 AGCTCAATTAATGAGTATGTTCCCATAGTAACGCCAATAGGAGCTTTCATTGACGTCA 181
Db 151 AGCTCAATTAATGAGTATGTTCCCATAGTAACGCCAATAGGAGCTTTCATTGACGTCA 210
Qy 182 TGGGTGAGTATTTACGGTAACTGCCCA-TTGGCAGTACATCAAGTATCATATATGCCA 240
Db 211 TGGGTGAGTATTTACGGTAACTGCCCACTTGGCAGTACATCAAGTATCATATATGCCA 270
Qy 241 AGTACGCCCTTATAGCTAATGAGCGTAAATGG 276
Db 271 AGTACGCCCTTATAGCTAATGAGCGTAAATGG 306

RESULT 11
LOCUS 108105
DEFINITION Sequence 3 from Patent EP 0309237.
ACCESSION 108105
VERSION 108105.1 GI:589184
KEYWORDS
SOURCE Unknown.
ORGANISM Unknown.
REFERENCE 1 (bases 1 to 777)
AUTHORS Gorman,C.M.
TITLE A transient expression system for producing recombinant protein
JOURNAL Patent: EP 0309237-A1 3 29-MAR-1989;
FEATURES location/Qualifiers
source 1..777
BASE COUNT 188 a 205 c 190 g 194 t
ORIGIN

Query Match 46.1%; Score 256; DB 6; Length 777;
Best Local Similarity 97.8%; Pred. No. 7e-41;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

Db 2 AGTATTATATAGTATCAATTAACGGGGTCATTAGTTCATATAGAGTTCCGC 61
Db 31 AGTATTATATAGTATCAATTAACGGGGTCATTAGTTCATATATAGAGTTCCGC 90
Qy 62 GTTACATTAACCTTACGTAATTTGGCCCGCGCTGACGCCCAAGAGCCCGCCCATTTG 121
Db 91 GTTACATTAACCTTACGTAATTTGGCCCGCGCTGACGCCCAAGAGCCCGCCCATTTG 150
Qy 122 AGCTCAATTAATGAGTATGTTCCCATAGTAACGCCAATAGGAGCTTTCATTGACGTCA 181
Db 151 AGCTCAATTAATGAGTATGTTCCCATAGTAACGCCAATAGGAGCTTTCATTGACGTCA 210

Qy 182 TGGGTGAGTATTTACGGTAACTGCCCA-TTGGCAGTACATCAAGTATCATATAGCCA 240
Db 211 TGGGTGAGTATTTACGGTAACTGCCCACTTGGCAGTACATCAAGTATCATATAGCCA 270
Qy 241 AGTACGCCCTTATAGCTAATGAGCGTAAATGG 276
Db 271 AGTACGCCCTTATAGCTAATGAGCGTAAATGG 306

RESULT 12
LOCUS BD000143
DEFINITION Vector having stabilized sequence and eucaryotic host cell.
ACCESSION BD000143
VERSION BD000143.1 GI:18623222
KEYWORDS JP 2000308497-A/1.
SOURCE unidentified.
ORGANISM unidentified.
REFERENCE 1 (bases 1 to 919)
AUTHORS Gorman,C.M.
TITLE Vector having stabilized sequence and eucaryotic host cell
JOURNAL Patent: JP 2000308497-A 1 07-NOV-2000;
COMMENT GENETIC INC
OS Unidentified
PN JP 2000308497-A/1
PD 07-NOV-2000
PF 17-APR-2000 JP 2000115248
PR 12-SEP-1986 US 907185,09-JUL-1987 US 071674 PI
CORNELIA MAXIN GOMAN
PC C12N15/09,C12N5/10,C12N15/00,C12N5/00
CC
FH key
FT source
FT 1..919
FEATURES location/Qualifiers
source 1..919
BASE COUNT 219 a 245 c 222 g 233 t
ORIGIN

Query Match 46.1%; Score 256; DB 6; Length 919;
Best Local Similarity 97.8%; Pred. No. 7e-41;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

Qy 2 AGTATTATATAGTATCAATTAACGGGGTCATTAGTTCATATAGAGTTCCGC 61
Db 31 AGTATTATATAGTATCAATTAACGGGGTCATTAGTTCATATATAGAGTTCCGC 90
Qy 62 GTTACATTAACCTTACGTAATTTGGCCCGCGCTGACGCCCAAGAGCCCGCCCATTTG 121
Db 91 GTTACATTAACCTTACGTAATTTGGCCCGCGCTGACGCCCAAGAGCCCGCCCATTTG 150
Qy 122 AGCTCAATTAATGAGTATGTTCCCATAGTAACGCCAATAGGAGCTTTCATTGACGTCA 181
Db 151 AGCTCAATTAATGAGTATGTTCCCATAGTAACGCCAATAGGAGCTTTCATTGACGTCA 210
Qy 182 TGGGTGAGTATTTACGGTAACTGCCCA-TTGGCAGTACATCAAGTATCATATAGCCA 240
Db 211 TGGGTGAGTATTTACGGTAACTGCCCACTTGGCAGTACATCAAGTATCATATAGCCA 270
Qy 241 AGTACGCCCTTATAGCTAATGAGCGTAAATGG 276
Db 271 AGTACGCCCTTATAGCTAATGAGCGTAAATGG 306

RESULT 13
LOCUS 105393
DEFINITION Sequence 14 from Patent EP 0260148.
ACCESSION 105393

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VERSION      105393.1  GI:591073
KEYWORDS
SOURCE       Unknown.
ORGANISM     Unknown.
REFERENCE    1 (bases 1 to 919)
AUTHORS      Gorman,C.M.
TITLE        Improved recombinant expression method, vector and transformed
              cells
JOURNAL      Patent: EP 0260148-A2 14 16-MAR-1988;
FEATURES     Location/Qualifiers
              source          1..919
              /organism="unknown"
BASE COUNT   219 a 245 c 222 g 233 t
ORIGIN
Query Match 46.1%; Score 256; DB 6; Length 919;
Best Local Similarity 97.8%; Pred. No. 7e-41;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 2 AGTATTAAATAGTAATCAATTAACGGGGTCAATTAGTTCATAGCCCATATATGAGATTCCGC 61
DB 31 AGTATTAAATAGTAATCAATTAACGGGGTCAATTAGTTCATAGCCCATATATGAGATTCCGC 90
QY 62 GTTACATAACTTACGGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 121
DB 91 GTTACATAACTTACGGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 150
QY 122 ACGTCATAATGACGTATGTTCCCATAGTAACGCCCATATGAGGACTTTCATGACGTCAA 181
DB 151 ACGTCATAATGACGTATGTTCCCATAGTAACGCCCATATGAGGACTTTCATGACGTCAA 210
QY 182 TGGGTGAGATTTTACGTAATCTGCAACTGCCCA-TTGGCAGTACATCAAGTATCATATGCCA 240
DB 211 TGGGTGAGATTTTACGTAATCTGCAACTGCCCACTTGGCAGTACATCAAGTATCATATGCCA 270
QY 241 AGTACGCCCTTATGACGTCAATGACGTAATATGG 276
DB 271 AGTACGCCCTTATGACGTCAATGACGTAATATGG 306

RESULT 14
LOCUS       108103          919 bp      DNA      linear      PAT 02-DEC-1994
DEFINITION  Sequence 1 from Patent EP 0309237.
ACCESSION   108103
VERSION     108103.1  GI:589182
KEYWORDS
SOURCE      Unknown.
ORGANISM    Unknown.
REFERENCE    1 (bases 1 to 919)
AUTHORS      Gorman,C.M.
TITLE        A transient expression system for producing recombinant protein
JOURNAL      Patent: EP 0309237-A1 1 29-MAR-1989;
FEATURES     Location/Qualifiers
              source          1..919
              /organism="unknown"
BASE COUNT   219 a 245 c 222 g 233 t
ORIGIN
Query Match 46.1%; Score 256; DB 6; Length 919;
Best Local Similarity 97.8%; Pred. No. 7e-41;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 2 AGTATTAAATAGTAATCAATTAACGGGGTCAATTAGTTCATAGCCCATATATGAGATTCCGC 61
DB 31 AGTATTAAATAGTAATCAATTAACGGGGTCAATTAGTTCATAGCCCATATATGAGATTCCGC 90
QY 62 GTTACATAACTTACGGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 121
DB 91 GTTACATAACTTACGGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 150

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QY 122 ACGTCATAATGACGTATGTTCCCATAGTAACGCCCATATGAGGACTTTCATGACGTCAA 181
DB 151 ACGTCATAATGACGTATGTTCCCATAGTAACGCCCATATGAGGACTTTCATGACGTCAA 210
QY 182 TGGGTGAGATTTTACGTAATCTGCAACTGCCCA-TTGGCAGTACATCAAGTATCATATGCCA 240
DB 211 TGGGTGAGATTTTACGTAATCTGCAACTGCCCACTTGGCAGTACATCAAGTATCATATGCCA 270
QY 241 AGTACGCCCTTATGACGTCAATGACGTAATATGG 276
DB 271 AGTACGCCCTTATGACGTCAATGACGTAATATGG 306

RESULT 15
LOCUS       AR028792          930 bp      DNA      linear      PAT 29-SEP-1999
DEFINITION  Sequence 2 from patent US 5858784.
ACCESSION   AR028792
VERSION     AR028792.1  GI:5940765
KEYWORDS
SOURCE      Unknown.
ORGANISM    Unknown.
REFERENCE    1 (bases 1 to 930)
AUTHORS      Debs,R.James. and Zhu,N.
TITLE        Expression of cloned genes in the lung by aerosol- and
              liposome-based delivery
JOURNAL      Patent: US 5858784-A 2 12-JAN-1999;
FEATURES     Location/Qualifiers
              source          1..930
              /organism="unknown"
BASE COUNT   233 a 228 c 211 g 258 t
ORIGIN
Query Match 46.1%; Score 256; DB 6; Length 930;
Best Local Similarity 97.8%; Pred. No. 7e-41;
Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 2 AGTATTAAATAGTAATCAATTAACGGGGTCAATTAGTTCATAGCCCATATATGAGATTCCGC 61
DB 157 AGTATTAAATAGTAATCAATTAACGGGGTCAATTAGTTCATAGCCCATATATGAGATTCCGC 216
QY 62 GTTACATAACTTACGGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 121
DB 217 GTTACATAACTTACGGTAATTTGGCCCGCGGTGACCGCCCAAGACCCCGCCCATTTG 276
QY 122 ACGTCATAATGACGTATGTTCCCATAGTAACGCCCATATGAGGACTTTCATGACGTCAA 181
DB 277 ACGTCATAATGACGTATGTTCCCATAGTAACGCCCATATGAGGACTTTCATGACGTCAA 336
QY 182 TGGGTGAGATTTTACGTAATCTGCAACTGCCCA-TTGGCAGTACATCAAGTATCATATGCCA 240
DB 337 TGGGTGAGATTTTACGTAATCTGCAACTGCCCACTTGGCAGTACATCAAGTATCATATGCCA 396
QY 241 AGTACGCCCTTATGACGTCAATGACGTAATATGG 276
DB 397 AGTACGCCCTTATGACGTCAATGACGTAATATGG 432

Search completed: December 15, 2002, 06:47:24
Job time : 3138 secs

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FE	XX	22-JUN-1999;	99JP-0174804.
PR	XX	22-JUN-1999;	99JP-0174804.
PA	XX	(JAPG) NIPPON ZEON KK.	
DR	XX	WPI; 2001-285232/30.	
PT	XX	New DNA molecule for use as a promoter for preparing a recombinant	
FT	XX	containing the DNA which is used for preparing a vaccine -	
PS	XX	Example 1; Page 11; 15pp; Japanese.	
CC	XX	The invention relates to a 5' and a 3' fragment of the chicken beta-actin	
CC	XX	promoter (AA84452 and AA84453, respectively), which exhibit promoter	
CC	XX	activity. The chicken beta-actin promoter fragments may be used to drive	
CC	XX	expression of a heterologous gene in a recombinant vector based as the	
CC	XX	active component of a genetic vaccine. The present sequence represents	
CC	XX	a chimeric promoter, pec, which was constructed in an exemplification of	
CC	XX	the invention. The pec promoter comprises a cytomegalovirus (CMV)	
CC	XX	immediate-early (IE) promoter fragment, and a 5' fragment of the chicken	
CC	XX	beta-actin promoter.	
SQ	XX	Sequence 561 BP; 109 A; 156 C; 191 G; 105 T; 0 other;	
Query Match		95.5%; Score 530; DB 22; Length 561;	
Best Local Similarity		99.5%; Pred. No. 3,2e-99;	
Matches	553;	Conservative 0; Mismatches 0; Indels 3; Gaps 2	
OY	2	AGTTAATAATGAATCAATTACGGGGCTCATTAGTTCATAGGCCCATATATGTGAGTTCGC 61	
DB	1	AGTTAATAATGAATCAATTACGGGGCTCATTAGTTCATAGGCCCATATATGTGAGTTCGC 60	
OY	62	GTTACAATACCTAGGTAATATGGCGCGGCTGACGCGCCCAAGAGACCCCGGCATTTG 121	
DB	61	GTTACAATACCTAGGTAATATGGCGCGGCTGACGCGCCCAAGAGACCCCGGCATTTG 119	
OY	122	ACGTCAATTAATGACGTATGTTCCCATAGTAGACCAATAGGACTTTCATTGACGTCAA 181	
DB	120	ACGTCAATTAATGACGTATGTTCCCATAGTAGACCAATAGGACTTTCATTGACGTCAA 179	
OY	182	TGGGTGGAGTATTACGGTAACATGCCCCATTGGCAGTACATCAAGTGATCATATGCCAA 241	
DB	180	TGGGTGGAGTATTACGGTAACATGCCCCATTGGCAGTACATCAAGTGATCATATGCCAA 239	
OY	242	GTACGCCCCCTTTTACGTCATGACGTAATATGATGAGTATTTTGTGCACGATGGG 301	
DB	240	GTACGCCCCCTTTTACGTCATGACGTAATATGATGAGTATTTTGTGCACGATGGG 299	
OY	302	GGC--GGGGGGGGGGGGCGCGCCAGCGCGGGGGGGGGGGGGGCGAGGGGCGGGG 359	
DB	300	GGC-----GGGGGGGGGGGGCGCGCCAGCGGGGGGGGGGGGGGCGAGGGGCGGGG 359	
OY	360	CGAGCGGAGAGAGTGTGGGGGCGAGCAATCAGAGCGCGCGCTCCGAAAAGTTTCCTTTA 419	
DB	360	CGAGCGGAGAGAGTGTGGGGGCGAGCAATCAGAGCGCGCGCTCCGAAAAGTTTCCTTTA 419	
OY	420	TGGCGAGCGGGCGCGCGCGCGCTTATAAAAAGCAAGCGCGCGGGGCGAGATCG 479	
DB	420	TGGCGAGCGGGCGCGCGCGCGCTTATAAAAAGCAAGCGCGCGGGGCGAGATCG 479	
OY	480	CTGGCGGCTGCTTGGCCCCCGTGGCCCCGCTCGCGCGCGCTCGCGCGCGCGCGCGCG 539	
DB	480	CTGGCGGCTGCTTGGCCCCCGTGGCCCCGCTCGCGCGCGCTCGCGCGCGCGCGCGCG 539	
OY	540	TCTGACTGACCGCGTC 555	
DB	540	TCTGACTGACCGCGTC 555	
RESULT 2			
ID	AAZ45930	standard; DNA; 6714 BP.	

XX	AA245930;
AC	
XX	
DT	25-APR-2000 (first entry)
XX	
DE	Nucleotide sequence of the vector p43CB-AT.
XX	
KM	Viral vector: p43CB-AT; alpha-1-antitrypsin; gene therapy; diabetes;
KM	alpha-1-antitrypsin deficiency; hemophilia; neurological disorder;
KW	adenosine deaminase deficiency; autoimmune disease; interleukin-2;
XX	immunodeficiency disease; infection; cytokine; circular; cancer; ss.
XX	
OS	Synthetic.
OS	Adeno associated virus.
OS	Homo sapiens.
OS	Cauliflower mosaic virus.
XX	Gallus sp.
PN	MO9955564-AI.
PD	04-NOV-1999.
XX	
PF	23-APR-1999; 99WO-US08921.
PR	24-APR-1998; 98US-0083025.
PA	(UYFL.) UNIV FLORIDA.
XX	
PI	Floette TR, Song S, Byrnes BJ, Morgan M;
DR	WPI; 2000-147020/13.
XX	
PT	Recombinant viral vector useful in the gene therapy of
PS	alpha-1-antitrypsin deficiency and also in hemophilia and diabetes -
XX	Claim 32; Fig 20A-C; 85pp; English.
CC	The present sequence represents a recombinant viral vector, designated
CC	p43CB-AT, which encodes human alpha-1-antitrypsin protein, and which
CC	contains the CMV enhancer, and chicken beta-actin promoter sequences.
CC	The vector is exemplary of the vectors of the invention which comprise
CC	a polynucleotide encoding a protein, and are used for gene therapy to
CC	correct genetic disorders related to expression of a protein of interest.
CC	The vectors of the invention are based on Adeno associated virus (AAV).
CC	The vectors comprise AAV inverted terminal repeats and constitutive or
CC	regulatable promoters for driving high levels of gene expression.
CC	Vectors encoding alpha-1-antitrypsin protein or a biologically active
CC	fragment or variant are administered to mammalian cells (preferably
CC	myofibers, myoblasts, hepatocytes or lung cells) for treating
CC	alpha-1-antitrypsin deficiency or ameliorating a condition resulting
CC	from a defective protein. The vectors can also be useful for genetic
CC	therapy of other conditions such as hemophilia, adenosine deaminase
CC	deficiency, diabetes, cancer, autoimmune diseases, neurological
CC	disorders, immunodeficiency diseases and bacterial or viral infections
CC	by the infusion of protein or a cytokine such as interleukin-2.
XX	
SO	Sequence 6714 BP; 1452 A; 1821 C; 1886 G; 1555 T; 0 other:
Query Match	58.6%; Score 325; DB 21; Length 6714;
Best Local Similarity	74.9%; Pred. No. 2e-57;
Matches	543; Conservative 0; Mismatches 10; Indels 172; Gaps
OY	2 AGTTTATTAGTATCAATTACGGGCGTCATTAGTTCATAGGCCCATATATGGAGTTCGC 61
DB	
323	AGTTTATTATAGTAATCAATTACGGGCGTCATTAGTTCATAGGCCCATATATGGAGTTCGC 382
OY	62 GTTACATACTAGCAGTAAATGGCCCCCGGTACCAGGCCAACAGACCCCAGCCATTG 121
DB	
383	GTTACATACTTAGTGCTAAATAGCCCCGCTGCTACCCGCCAACAGACCCCAGCCATTG 442
OY	122 ACGTCAATATGAGCTATGTTCCTCATAGTACGCCAATATAGGACTTTCATTGACGTCAA 181
DB	
443	ACGTCAATATGAGCTATGTTCCTCATAGTACGCCAATATAGGACTTTCATTGACGTCAA 502

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OY 182 TGGGTGAGATATTACGGTAACTGCCCA-TTGGCAGTACATCAAGTGTATCATATGCCA 240
    |||||||
DB 503 TGGGTGAGATATTACGGTAACTGCCCA-TTGGCAGTACATCAAGTGTATCATATGCCA 562
OY 241 AGTACGCCCCCTATTGACGTCAATGACGTTAAATG----- 276
    |||
DB 563 AGTCCGCCCTATTGACGTCAATGACGTTAAATGAGCCCTGCGCATTTATGCCAGTAC 622
OY 277 ----- 276
DB 623 ATGACCTTACGGACTTCTCACTTGCGAGTACATCTACATCTATGCTATTCCTATTAC 682
OY 277 ----- 276
DB 683 ATGTCGAGTGTAGAGCCCACTTCCTCACTTCCCATCTCCCTCCCTCCACCC 742
OY 277 -----ATGCACTATTGTCAGCAGTGGGGC-----GGGGGG 310
    |||
DB 743 CCAATTTTGTATTATTATTATTATTATTATTATTATTATTGTCAGCATGAGGGGGGGGG 802
OY 311 GGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGAGA 370
    |||||||
DB 803 GGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGAGA 862
OY 371 GGTGCGGCGGAGCCATCAGACGCGCGCTCCGAAAGTTTCCTTTATGCGAGGCGG 430
    |||||||
DB 863 GGTGCGGCGGAGCCATCAGACGCGCGCTCCGAAAGTTTCCTTTATGCGAGGCGG 922
OY 431 GGTGCGGCGGCGCTTATTAAGCGAAAGCGCGGGGGGGGGAGTCCGTCGCG-CGCTG 489
    |||||||
DB 923 CGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGTGG 982
OY 490 CTTTCGCGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG 549
    |||||||
DB 983 CTTTCGCGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG 1042
OY 550 CGCGT 554
    |||||
DB 1043 CGCGT 1047

```

RESULT 3

AAZ45934
ID AAZ45934 standard; DNA; 6924 BP.

AAZ45934:

25-APR-2000 (first entry)

Nucleotide sequence of the vector p43msENCB-AT.

Viral vector: p43msENCB-AT; alpha-1-antitrypsin; gene therapy; diabetes;

alpha-1-antitrypsin deficiency; haemophilia; neurological disorder;

adenosine deaminase deficiency; autoimmune disease; interleukin-2;

immunodeficiency disease; infection; cytokine; circular; cancer; ss.

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xx

xx

PI Plote TR, Song S, Byrnes BJ, Morgan M;
XX WPI; 2000-147020/13.
XX
XX
PT Recombinant viral vector useful in the gene therapy of
PT alpha-1-antitrypsin deficiency and also in hemophilia and diabetes -
PS Claim 32; Fig 24A-C; 85pp; English.
XX
XX
CC The present sequence represents a recombinant viral vector, designated
CC p43msENCB-AT, which encodes human alpha-1-antitrypsin protein, and
CC which contains the CMV enhancer, and chicken beta actin promoter
CC sequences, and has an enhancer sequence upstream of the CMV promoter.
CC The vector is exemplary of the vectors of the invention which comprise
CC a polynucleotide encoding a protein, and are used for gene therapy to
CC correct genetic disorders related to expression of a protein of interest.
CC The vectors of the invention are based on Adeno associated virus (AAV).
CC The vectors comprise AAV inverted terminal repeats and constitutive or
CC regulatable promoters for driving high levels of gene expression.
CC Vectors encoding alpha-1-antitrypsin protein or a biologically active
CC fragment or variant are administered to mammalian cells (preferably
CC myofibers, myoblasts, hepatocytes or lung cells) for treating
CC alpha-1-antitrypsin deficiency or ameliorating a condition resulting
CC from a defective protein. The vectors can also be useful for genetic
CC therapy of other conditions such as hemophilia, adenosine deaminase
CC deficiency, diabetes, cancer, autoimmune diseases, neurological
CC disorders, immunodeficiency diseases and bacterial or viral infections
CC by the infusion of protein or a cytokine such as interleukin-2.
XX
SQ Sequence 6924 BP; 1504 A; 1866 C; 1951 G; 1603 T; 0 other;

Query Match 58.6%; Score 325; DB 21; Length 6924;
Best Local Similarity 74.9%; Pred. No. 2e-57;
Matches 543; Conservative 0; Mismatches 10; Indels 172; Gaps 4;

```

OY 2 AGTTATTAATAGTATCAATTTACGGGCTCATTTGTTTCATATGCCCCATATATGAGTTCCG 61
    |||||||
DB 533 AGTTATTAATAGTATCAATTTACGGGCTCATTTGTTTCATATGCCCCATATATGAGTTCCG 592
OY 62 GTTACATTAATAGTATCAATTTACGGGCTCATTTGTTTCATATGCCCCATATATGAGTTCCG 121
    |||||||
DB 593 GTTACATTAATAGTATCAATTTACGGGCTCATTTGTTTCATATGCCCCATATATGAGTTCCG 652
OY 122 ACSTCAATATGACGATGTTTCCCATATGTAACCCATATGAGATTTCCATGACGTCAA 181
    |||||||
DB 653 ACSTCAATATGACGATGTTTCCCATATGTAACCCATATGAGATTTCCATGACGTCAA 712
OY 182 TGGGTGAGATATTACGTAACCTGCCA-TTGGCAGTACATCAAGTGTATCATATGCCA 240
    |||||||
DB 713 TGGGTGAGATATTACGTAACCTGCCA-TTGGCAGTACATCAAGTGTATCATATGCCA 772
OY 241 AGTACGCCCCCTATTGACGTCAATGACGTTAAATG----- 276
    |||
DB 773 AGTCCGCCCTATTGACGTCAATGACGTTAAATGAGCCCTGCGCATTTATGCCAGTAC 832
OY 277 ----- 276
DB 833 ATGACCTTACGGACTTCTCACTTGCGAGTACATCTACGTTATGTCATCGCTATTAC 892
OY 277 ----- 276
DB 893 ATGTCGAGTGTAGAGCCCACTTCCTCACTTCCCATCTCCCTCCCTCCACCC 952
OY 277 -----ATGCACTATTGTCAGCAGTGGGGC-----GGGGGG 310
    |||
DB 953 CCAATTTTGTATTATTATTATTATTATTATTATTATTATTGTCAGCATGAGGGGGGGGG 1012
OY 311 GGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGAGA 370
    |||||||
DB 1013 GGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGAGA 1072
OY 371 GGTGCGGCGGAGCCATCAGACGCGCGCTCCGAAAGTTTCCTTTATGCGAGGCGG 430
    |||||||

```

Db	1073	GGTGGGCGGACAGCATGACGAGGGGCGGCTCCGAAATTTCTTTATGGCGAGGGG	1132
Oy	431	CGGCGGCGGCGCCCTATATAAAGCGAGCGCGCGGAGGAGTGGCTGG-CGCTG	489
Db	1133	CGGCGGCGGCGGCGCTATATAAAGCGAGCGCGGCGGAGGAGTGGCTGGCGAGCGTG	1192
Oy	490	CGTTTGGCGCGGTTGGCGCGCGCGCGCTTGGCGCGCGCGCGCGCGCGCTGTGACTGAC	549
Db	1193	CGTTTGGCGCGGTTGGCGCGCGCGCGCTTGGCGCGCGCGCGCGCGCGCTGTGACTGAC	1252
Oy	550	CGCGT 554	
Db	1253	CGCGT 1257	
RESULT 4			
ID	AAZ45935		
AC	AAZ45935 standard; DNA: 6924 BP.		
XX	AAZ45935;		
DT	25-APR-2000 (first entry)		
XX			
DE	Nucleotide sequence of the vector p43msENCB-AT.		
XX			
KM	Viral vector: p43msENCB-AT; alpha-1-antitrypsin; gene therapy; diabetes;		
KM	alpha-1-antitrypsin deficiency; haemophilia; neurological disorder;		
KM	adenosine deaminase deficiency; autoimmune disease; interleukin-2;		
OS	immunodeficiency disease; infection; cytokine; circular; cancer; ss.		
XX			
OS	Synthetic.		
OS	Adeno associated virus.		
OS	Homo sapiens.		
OS	Cauliflower mosaic virus.		
OS	Gallus sp.		
PN	MO995564-A1.		
PD			
XX	04-NOV-1999.		
XX			
PF	23-APR-1999; 99MO-US08921.		
XX			
PR	24-APR-1998; 98US-0083025.		
XX			
PA	(UYFL) UNIV FLORIDA.		
XX			
PI	Flotte TR, Song S, Byrnes BJ, Morgan M;		
XX			
DR	WPI: 2000-147020/13.		
XX			
PT	Recombinant viral vector useful in the gene therapy of		
XX	alpha-1-antitrypsin deficiency and also in hemophilia and diabetes -		
PS	Claim 32; Fig 25A-C; 85pp: English.		
CC	The present sequence represents a recombinant viral vector, designated		
CC	p43msENCB-AT, which encodes human alpha-1-antitrypsin protein, and		
CC	which contains the CMV enhancer, and chicken beta-actin promoter		
CC	sequences, and has an enhancer sequence upstream of the CMV promoter		
CC	in the opposite direction to the enhancer of vector p43msENCB-AT.		
CC	The vector is exemplary of the vectors of the invention which comprise		
CC	a polynucleotide encoding a protein, and are used for gene therapy to		
CC	correct genetic disorders related to expression of a protein of interest		
CC	The vectors of the invention are based on Adeno associated virus (AAV).		
CC	The vectors comprise AAV inverted terminal repeats and constitutive or		
CC	regulatable promoters for driving high levels of gene expression.		
CC	Vectors encoding alpha-1-antitrypsin protein or a biologically active		
CC	fragment or variant are administered to mammalian cells (preferably		
CC	myoblasts, myoblasts, hepatocytes or lung cells) for treating		
CC	alpha-1-antitrypsin deficiency or ameliorating a condition resulting		
CC	from a defective protein. The vectors can also be useful for genetic		
CC	therapy of other conditions such as hemophilia, adenosine deaminase		
CC	deficiency, diabetes, cancer, autoimmune diseases, neurological		

CC	disorders; immunodeficiency diseases and bacterial or viral infections
CC	by the infusion of protein or a cytokine such as interleukin-2.
XX	
SO	Sequence 6924 BP; 1500 A; 1886 C; 1931 G; 1607 T; 0 other:
	Query Match 58.6%; Score 325; DB 21; Length 6924;
	Best Local Similarity 74.9%; Pred. No. 2e-57;
	Matches 543; Conservative 0; Mismatches 10; Indels 172; Gaps 4;
OY	2 AGTTATTAATAAGTAACTACGCGGGTCATTAGTTCATAGGCCAATATATGAGTTCCGC 61 DB 533 AGTATTAATAGTAAATCAATACGCGGGTCATTAGTTCATAGGCCAATATATGAGTTCCGC 592
OY	62 GTTACATTAACCTACGATAATGTGGCCCCCGGCTGACGCGCCCAACAGACCOCGCCATTG 121 DB 593 GTTACATTAACCTACGATAATGTGGCCCCCGGCTGACGCGCCCAACAGACCOCGCCATTG 652
OY	122 ACGCAATAATAGCATGTGTTCCCATAGTAAGGCCAATATGAGACTTTCATTGAGCTCA 181 DB 653 ACGCAATAATAGCATGTGTTCCCATAGTAAGGCCAATATGAGACTTTCATTGAGCTCA 712
OY	182 TGGGAGGAGTATTTACGGAATACGCGCA-TGGCAGTAGTCAATCAAGTATCATATGCA 240 DB 713 TGGGAGGAGTATTTACGGAATACGCGCACTTGCCACTGGCAGTACATCAAGTATCATATGCA 772
OY	241 AGTACGCCCCCTATTGACGTCAATGACGTAATG----- 276 DB 773 AGTACGCCCCCTATTGACGTCAATGACGTAATGACGCGCCCTGCGCATTAATGCCCAGTAC 832
OY	277 ----- 276
Db	833 ATGACCTTAACGGGACTTTCCTACTTGCGAGTACATCTAGTATGATCATGCTATTAC 892
OY	277 ----- 276
Db	893 ATGTCGAGTGAAGCCCAAGCTTCCTACTTGCGAGTACATCTAGTATGATCATGCTATTAC 952
OY	277 -----ATGCAGTATTTTGTGCAGCGATGGGGC---GGGGGG 310 DB 953 CCAATTTTGTATTATTATTATTTTAAATATTTTGTGCAGCATGGGGCGGGGGGGGG 1012
OY	311 GGGGGGGCGCGCCACAGCGGGGGCGGGCGAGGGCGGGGGCGGGCGAGGGGAGA 370 DB 1013 GGGGGGGCGCGCCACAGCGGGGGCGGGCGAGGGCGGGGGCGGGCGAGGGGAGA 1072
OY	371 GGTGGCGCGGCGCAATACAGAGCGGCGGCTCCGAAATTTCTTTTATGCGGAGCGG 430 DB 1073 GGTGGCGCGGCGCAATACAGAGCGGCGGCTCCGAAATTTCTTTTATGCGGAGCGG 1132
OY	431 CGGCGGCGGCGCCCTATATAAAAAGCGAAGCGCGGGCGGGGAGATCGCTGCG-CGCTG 489 DB 1133 CGGCGGCGGCGCCCTATATAAAAAGCGAAGCGCGGGCGGGGAGATCGCTCGAGCGTG 1192
OY	490 CCTTGGCCCGCTGCGCGCGCTCGCGCGGCGCTCGCGCGCGCCCGCGGCTGTGACTAC 549 DB 1193 CCTTGGCCCGCTGCGCGCGCTCGCGCGGCGCTCGCGCGCGCCCGCGGCTGTGACTAC 1252
OY	550 CGCGT 554
Db	1253 CGCGT 1257
RESULT 5	
AAH21793	
ID	AAH21793 standard; cDNA; 4211 BP.
XX	AAH21793;
XX	15-AUG-2001 (first entry)
DT	Mouse Megl/grb10 nucleotide sequence SEQ ID NO:3.
DE	
XX	Mouse; Megl/grb10; diabetes; transgene; transgenic animal;

XX	Insulin signal transduction inhibition; ss.
XX	OS Mus sp.
XX	Key Location/Qualifiers
XX	Key CDS 1862..3652
XX	FT /tag= a
XX	FT /product= "Meg1/Grb10"
XX	WT0200128321-A1.
XX	26-APR-2001.
XX	18-AUG-2000; 2000WO-JP0546.
XX	20-OCT-1999; 99JP-0298273.
XX	(NISC-) JAPAN SCI & TECHNOLOGY CORP.
XX	Ishino F, Miyoshi N, Ishino T, Yokoyama M, Wakana S;
XX	WPI: 2001-300253/31.
XX	P-PSDB: AAB98059.
XX	Transgenic non-human mammal with Meg1/Grb10 or human GRB 10 gene useful as a model for onset of diabetes and for screening new diabetes treatments
XX	Claim 5; Page 32-33; 50pp; Japanese.
XX	The present invention describes a transgenic non-human mammal containing the Meg1/Grb10 gene. Also described are: (1) a transgenic non-human mammal with human GRB10 gene; (2) a method for producing a transgenic mouse; (3) method (M1) for screening for drugs for treating diabetes; and (4) drugs found using (M1). The transgenic non-human mammal is useful for screening for new drugs to treat diabetes. The transgenic animals are models for the onset of diabetes, and may be useful in discovering the mechanism for the onset of diabetes caused by inhibition of Insulin signal transduction, and for developing new treatments. The present sequence represents a specifically claimed transgene comprising the mouse Meg1/Grb10 nucleotide sequence downstream of the chicken beta-actin promoter and upstream of rabbit beta-globin polyA.
XX	Sequence 4211 BP; 890 A; 1139 C; 1230 G; 952 T; 0 other;
XX	Query Match 55.78; Score 309.4; DB 22; Length 4211;
XX	Best Local Similarity 74.68; Pred. No. 2.9e-54;
XX	Matches 539; Conservative 0; Mismatches 11; Indels 173; Gaps 5;
YY	2 AGTTATTAATGATCAATTAACGGGGTCATTAGTCCATATATGAGTTCGGC 61
YY	
YY	138 AGTATTAATTAATCAATTAACGGGGTCATTAGTCCATATATGAGTTCGGC 197
YY	62 GTTACATACTTACGGTAAATTTGGCCGGCCGTACCGCCCAAGACCCCGCCCATTTG 121
YY	198 GTTACATACTTACGGTAAATTTGGCCGGCCGTACCGCCCAAGACCCCGCCCATTTG 257
YY	122 ACGTTAATTAATGACGTATATGTTCCCATATGTAAGACCAATATGAGCATTTCCATGAGCTCA 181
YY	258 ACGTTAATTAATGACGTATATGTTCCCATATGTAAGACCAATATGAGCATTTCCATGAGCTCA 317
YY	182 TGGGTGAGATATTTACGGTAAACTCCCA-TTGGCAGTACATCAAGTATCATATGCCA 240
YY	318 TGGGTGAGATATTTACGGTAAACTCCCACTTGGCAGTACATCAAGTATCATATGCCA 377
YY	241 AGTACGCCCTTATGACGTCAATGACGTTAAATGG----- 276
YY	378 AGTACGCCCTTATGACGTCAATGACGTTAAATGGCCGCTGACATTAATGCCAGTAC 437
YY	277 ----- 276
YY	438 ATGACCTTAATGAGGACTTTCCTACTTGGCAGTACATCAATCAATTAATGATCGCATTTACC 497

Oy	277	-----	276
Db	48	ATGGGTCAAGGAGAGCCCAAGCTTCTGCTACCTCCCAATCTCCCCCTCCCAAC	557
Oy	277	-----ATGCAATATTTTGTGCAGCATGAGGAGC--GGGGGG	311
Db	558	CCCAATTTTGTATTTATTTATTTTAAATTTTGTGACAGATGGGGGGGGGG	617
Oy	312	GGGGGCGCGCCCAAGCGGGGCGGAGGGGCGGGGCGGAGCGAAG	371
Db	618	GGGGGCGCGCGCAAGCGGGGCGGAGGGGCGGGGCGGAGCGAAG	677
Oy	372	GTGGCGGGGCAAGCAATAGAGCGGCGTCCGAAAGTTTCTTTATAGGAGGCG	431
Db	678	GTGGCGGGGCAAGCAATAGAGCGGCGTCCGAAAGTTTCTTTATAGGAGGCG	737
Oy	432	GGCGCGCGCGCCTATATAAAGCAAGCGCGCGGCGGAGTCCGCGCTCC	491
Db	738	GGCGCGCGCGCCTATATAAAGCAAGCGCGCGGCGGAGTCCCT--GCCTGCC	795
Oy	492	TTTGGCCCGTGGCCGCTCCGCGCGCGCTCGCGCGCGCGCGCGCTGACTGACG	551
Db	796	TTTGGCCCGTGGCCGCTCCGCGCGCGCTCGCGCGCGCGCGCGCTGACTGACG	854
Oy	552	CGT 554	
Db	855	CGT 857	
RESULT 6			
AAL43172			
ID	AAL43172	standard; DNA; 5504 BP.	
XX	AAL43172;		
DE	16-AUG-2002	(first entry)	
DE	Rat expression product activity regulation method-related DNA sequence.		
XX	Norway rat; ds; expression product activity regulation.		
OS	Rattus norvegicus.		
PN	WO200241922-A1.		
XX	30-MAY-2002.		
PF	28-SEP-2001; 2001WO-JP08575.		
XX	24-NOV-2000; 2000JP-0358389.		
PA	(CHUS) CHUGAI SEIYAKU KK.		
PA	(MARU/) MARUYAMA H.		
PI	Maruyama H, Miyazaki J, Sugawa M, Higuchi M;		
DR	WPI: 2002-471702/50.		
PT	Regulating activity of expression product of gene transferred into living body by coexistence of interfering substance e.g. expression product receptor, given through skin to express and reduce excessive activity		
PS	Example 1; Page 35-39; 49pp; Japanese.		
CC	The invention comprises a method for regulating the activity of an expression product of a gene transferred into a living body. The method involves the coexistence of a protein interfering with the activity of the expression product. The method of the invention is useful for regulating the activity of an expression product. The present rat DNA sequence was used in an example of the invention.		
CC	Sequence 5504 BP; 1159 A; 1491 C; 1518 G; 1336 T; 0 other;		

Query Match	55.78;	Score 309.4;	DB 24;	Length 5504;
-------------	--------	--------------	--------	--------------

Best Local Similarity: 74.6%; Pred. No. 2.9e-54;
Matches 539; Conservative 0; Mismatches 11; Indels 173; Gaps 5.

	QY	2	AGTATTAAATGTGTAATCAATTAAGGGGTCTATTAGTTCATATGAGCCATATATGAGGTCCG	61
	Db	21	AGTTATTATATAGTATCAATTTACGGGGTCTATTAGTTCATATGAGCCATATATGAGGTCCG	80
QY	62	GTTCACATTAACCTACGTAATTTGGCCCGCGGTGACCGGCCAACGACCCCGCCATTG	121	
Db	81	GTTCACATTAACCTACGTAATTTGGCCCGCGGTGACCGGCCAACGACCCCGCCATTG	140	
QY	122	ACGCATTAATTAATGAGTATGTCTCCATATGATACCCCAATAGGACTTTCATATGAGCTCAA	181	
Db	141	ACGCATTAATTAATGAGTATGTCTCCATATGATACCCCAATAGGACTTTCATATGAGCTCAA	200	
QY	182	TGGGTGAGAGTATTAACAGTAACTGGCCA-TTGGCAGTACATCAAGTATCATATGACA	240	
Db	201	TGGGTGAGAGTATTAACAGTAACTGGCCACTTGGCAGTACATCAAGTATCATATGACA	260	
QY	241	AGTACGCCCCCTATTGACGTCATGACGGTAATGC-----	276	
Db	261	AGTACGCCCCCTATTGACGTCATGACGGTAATGC-----	320	
QY	277	-----	276	
Db	321	ATGACCTTATAGGACTTCTCTACTTGCGAGTACATCTACGTATTAAGTCATCCGTATTACC	380	
QY	277	-----	276	
Db	381	ATGGGTGAGGTGAGCCCCACGTTCTGCTCACTCTCCCATCTCCCCCCTCCACAC	440	
QY	277	-----ATGAGATTTTGTGACAGGATGGGGG--GGGGGGG	311	
Db	441	CCCAATTTTGTATTTATTTATTTTATTTATTTTGTGACAGGATGGGGG--GGGGGGG	500	
QY	312	GGGGGCGCGGCCAGGCGGGGCGGGGCGAGGCGCGGGCGGGCGAGGCGAGAG	371	
Db	501	GGGGGCGCGGCCAGGCGGGGCGGGGCGAGGCGGGCGGGCGAGGCGAGAG	560	
QY	372	GTCGGGGGGGAGCAATCAGAGGGCGGCTCGGAAAGTTCTTTATAGGAGGCGGC	431	
Db	561	GTCGGGGGGGAGCAATCAGAGGGCGGCTCGGAAAGTTCTTTATAGGAGGCGGC	620	
QY	432	GGCGGCGCGGCCCTATATAAAGCGAAGCGCGGGCGGGAGGATCGCTGCGCGCTGCC	491	
Db	621	GGCGGCGCGGCCCTATATAAAGCGAAGCGCGGGCGGGAGGATCGCT--CGCTGGC	678	
QY	492	TTTCGCCCGGTGCCCCGCTCGCGCGCGGCTCGCGCGGCCCGCGGGCTGTGACTGACCG	551	
Db	679	TTTCGCCCGGTGCCCCGCTCGCGCGCGGCTCGCGCGGCCCGCGGGCTGTGACTGACCG	737	
QY	552	CGT 554		
Db	738	CGT 740		
RESULT 7				
AAL43171				
ID AAL43171 standard; DNA: 5565 BP.				
XX AAL43171:				
AC	16-AUG-2002	(first entry)		
XX				
DE				
XX				
FW				
XX				
OS				
NN				
Human: ds; expression product activity regulation method-related DNA sequence				
Homo sapiens.				
W0200241922-A1.				

XX (DELT-) DELTAGEN INC.
 XX
 XX Siebel C, Brennan TJ;
 XX
 XX WPI: 2002-383132/41.
 XX
 PT Novel targeting vector modifying target gene, has first and second
 PT sequences homologous to target gene portions, a selectable marker
 PT cassette and regulator, useful for producing animals with targeted gene
 PT modifications
 XX
 XX Example 1: Fig 14A-B; 43pp: English.
 XX
 CC The invention describes a targeting vector (positive selection vector)
 CC (I) capable of modifying a target gene. (I) comprises two sequences
 CC (S1 and S2) homologous to a portion or region of a target gene, a
 CC selectable marker cassette and a regulator. (I) is useful for producing
 CC cells comprising a modification of the target gene which involves
 CC introducing (I) into cells capable of homologous recombination, selecting
 CC for cells expressing the selectable marker and identifying cells
 CC containing the modification or modification of target gene enhances
 CC cells comprising disruption or modification of target gene
 CC recovery of cells having targeting vector integrated via homologous
 CC recombination into the genomes of the cells. (I) is capable of modifying
 CC target gene in a cell with high efficiency and specificity. Use of (I)
 CC provides a faster and more efficient means for isolating and selecting
 CC cells comprising target gene modification. Also use of (I) provides an
 CC increase over previous technologies in both the speed and frequency at
 CC which homologous recombination events can be recovered. (I) is also
 CC useful for creation of transgenic animals containing targeted gene
 CC modifications. This sequence represents the positive selection vector
 CC construct c3408.
 CC
 XX Sequence 5759 BP; 1274 A; 1526 C; 1658 G; 1301 T; 0 other;
 XX
 SQ
 Query Match 55.7%; Score 309.4; DB 24; Length 5759;
 Best Local Similarity 74.6%; Pred. No. 2.9e-54;
 Matches 539; Conservative 0; Mismatches 11; Indels 173; Gaps 5;

DB 539 GGGGGCGCGCGCCAGAGCGGGGGCGGGCGGAGAGGGCGGGCGGAGCGGAGAG 598
 |||||||
 QY 372 GTGCGGCGGCGAGCCCAATGAGCGCGCGCTCCGAAAGTTCTTTATGAGCGCGC 431
 |||||||
 DB 599 GTGCGGCGGCGAGCCCAATGAGCGCGCGCTCCGAAAGTTCTTTATGAGCGCGC 658
 |||||||
 QY 432 GCGGCGGCGCGCTTTATTAAGCGAGCGCGCGCGGAGTGTGCGCGCTGCGC 491
 |||||||
 DB 659 GCGGCGGCGCGCTTTATTAAGCGAGCGCGCGCGGAGTGTGCT--GCGTTGCG 716
 |||||||
 QY 492 TTCGCGCGCGCTGCGCGCGCGCGCGCGCGCGCGCGCGCGCTGAGTGAACG 551
 |||||||
 DB 717 TTCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCTGAGTGAACG 775
 |||||||
 QY 552 CGT 554
 |||
 DB 776 CGT 778
 |||
 RESULT 10
 ID ABR49520 standard; DNA; 6148 BP.
 XX ABR49520
 AC ABR49520;
 XX
 DT 15-JUL-2002 (first entry)
 XX
 DE Positive selection vector construct c3406.
 XX
 KM Transgenic animal; targeting vector; positive selection vector;
 KM homologous recombination; target gene modification; transgenic animal;
 KM c3406; ds.
 XX
 OS Synthetic.
 XX
 PN WO200222834-A2.
 XX
 PD 21-MAR-2002.
 XX
 PF 17-SEP-2001; 2001WO-0528892.
 XX
 PR 15-SEP-2000; 2000US-232957P.
 XX
 PA (DELT-) DELTAGEN INC.
 XX
 PI Siebel C, Brennan TJ;
 XX
 WPI: 2002-383132/41.
 DR
 PT Novel targeting vector modifying target gene, has first and second
 PT sequences homologous to target gene portions, a selectable marker
 PT cassette and regulator, useful for producing animals with targeted gene
 PT modifications
 XX
 XX Claim 32; Fig 6A; 43pp: English.
 XX
 CC The invention describes a targeting vector (positive selection vector)
 CC (I) capable of modifying a target gene. (I) comprises two sequences
 CC (S1 and S2) homologous to a portion or region of a target gene, a
 CC selectable marker cassette and a regulator. (I) is useful for producing
 CC cells comprising a modification of the target gene which involves
 CC introducing (I) into cells capable of homologous recombination, selecting
 CC for cells expressing the selectable marker and identifying cells
 CC containing the modification of the target gene. Use of (I) for enriching
 CC cells comprising disruption or modification of target gene enhances
 CC recovery of cells having targeting vector integrated via homologous
 CC recombination into the genomes of the cells. (I) is capable of modifying
 CC target gene in a cell with high efficiency and specificity. Use of (I)
 CC provides a faster and more efficient means for isolating and selecting
 CC cells comprising target gene modification. Also use of (I) provides an
 CC increase over previous technologies in both the speed and frequency at
 CC which homologous recombination events can be recovered. (I) is also

Db 441 CCCAATTGTGTAATTATTTTAAATTATTTGTGCAGCGATGGGGCGGGGGGG 50

QY 312 GGGGGCGCCGCGCAGGCGGGGGCGGGGCGAGGCGGGCGGGCGAGCGAGAG 371
DB 501 GGGGGCGCCGCGCAGGCGGGGGCGGGGCGAGGCGGGGGCGGGCGAGAG 560
QY 372 GTGGCGCGGCGAGCAATCAGAGCGGCGCTCCGAAAGTTTCTTTATGCGAGCGGC 431
DB 561 GTGGCGCGGCGAGCAATCAGAGCGGCGCTCCGAAAGTTTCTTTATGCGAGCGGC 620
QY 432 GCGCGCGCGCGCCCTATATAAAGCGAAGCGCGCGCGGCGGAGTCCGCGCTGCC 491
DB 621 GCGCGCGCGCGCCCTATATAAAGCGAAGCGCGCGCGGCGGAGTCCGCT - GCGTTGCC 678
QY 492 TTGCGCGCGCGCTGCGCTCCGCGCGCGCTGCGCGCGCGCGCGCTGACTACCG 551
DB 679 TTGCGCGCGCGCTGCGCTCCG - GCGCGCGCTGCGCGCGCGCGCGCTGACTACCG 737
QY 552 CGT 554
DB 738 CGT 740

RESULT 12

AAFB4456

ID AAFB4456 standard; DNA: 599 BP.

AC AAFB4456;

DT 25-JUN-2001 (first entry)

DE Cytomegalovirus (CMV) immediate-early (IE) promoter enhancer.

XX Cytomegalovirus; CMV; immediate-early; IE promoter enhancer;

KM Pec promoter; chicken beta-actin promoter;

KW recombinant vector; genetic vaccine; gene therapy; ds.

XX Human cytomegalovirus.

OS JP2001000188-A.

PM 09-JAN-2001.

PD 22-JUN-1999; 99JP-0174804.

PF 22-JUN-1999; 99JP-0174804.

PR 22-JUN-1999; 99JP-0174804.

XX (JARG) NIPPON ZERON KK.

PA WPI: 2001-285232/30.

DR WPI: 2001-285232/30.

XX New DNA molecule for use as a promoter for preparing a recombinant

PT containing the DNA which is used for preparing a vaccine -

PT Example 1; Page 12; 15pp: Japanese.

PS The invention relates to a 5' and a 3' fragment of the chicken beta-actin

XX promoter (AAFB4452 and AAFB4453, respectively), which exhibit promoter

CC activity. The chicken beta-actin promoter fragments may be used to drive

CC expression of a heterologous gene in a recombinant vector used as the

CC active component of a genetic vaccine. The present sequence represents

CC a cytomegalovirus (CMV) immediate-early (IE) promoter enhancer, a

CC portion of which was used in the construction of the chimeric Pec

CC promoter (AAFB4454) in an exemplification of the invention.

XX Sequence 599 BP; 152 A; 144 C; 144 G; 159 T; 0 other;

QY Query Match 46.3%; Score 257; DB 22; Length 599;

Best Local Similarity 97.8%; Pred. No. 1.3e-43;

Matches 271; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 1 GAGTATTAATAGTAAATCAATACGGGGGCTAGTTCATAGCCCAATATGAGTTCCG 60

DB 6 GAGTATTAATAGTAAATCAATACGGGGGCTAGTTCATAGCCCAATATGAGTTCCG 65

QY 61 CGTTACATTAATAGTAAATTTGGCCCGCGGCTGACCGCCCAAGACCCCGGCCATT 120
DB 66 CGTTACATTAATAGTAAATTTGGCCCGCGGCTGACCGCCCAAGACCCCGGCCATT 125
QY 121 GAGCTAATTAATAGTAAATTTGGCCCGCGGCTGACCGCCCAAGACCCCGGCCATT 180
DB 126 GAGCTAATTAATAGTAAATTTGGCCCGCGGCTGACCGCCCAAGACCCCGGCCATT 185
QY 181 ATGGGTGAGTATTTACGGTAAACTGCCA - TTGGCAGTACATCAAGTATATATGCG 239
DB 186 ATGGGTGAGTATTTACGGTAAACTGCCA - TTGGCAGTACATCAAGTATATATGCG 245
QY 240 AAGTACGCGCCCTATTTAGCTCAATGAGTAAATG 276
DB 246 AAGTACGCGCCCTATTTAGCTCAATGAGTAAATG 282

RESULT 13

AAK15163/C

ID AAK15163 standard; DNA: 589 BP.

AC AAK15163;

DT 22-APR-1999 (first entry)

DE Human cytomegalovirus promoter region.

XX apolipoprotein; binding; in vivo transport; nucleic acid; binding domain;

KM nuclear localisation sequence; gene therapy; cancer; cystic fibrosis;

KW non-small cell lung carcinoma; diabetes; arteriosclerosis;

XX Human cytomegalovirus promoter region; ss.

OS Human cytomegalovirus.

PN WO9856938-A1.

PD 17-DEC-1998.

PF 10-JUN-1998; 98WO-0511927.

PR 14-MAY-1998; 98US-0079030.

PR 13-JUN-1997; 97US-0874807.

XX (BAYU) BAYLOR COLLEGE MEDICINE.

PI Guevara JG, Hoogeveen RC, Moore JP;

PA WPI: 1999-070331/06.

DR WPI: 1999-070331/06.

XX Composition comprising nucleic acid bound to LDL or VLDL lipoprotein

PT - used for delivering nucleic acid to cells for gene therapy and

PT antisense treatment

PS Example 4; Page 74; 293pp: English.

XX The present sequence represents the Human cytomegalovirus promoter

CC region. The compositions of the invention bind to this sequence.

CC The specification describes a composition that comprises low

CC density lipoprotein (LDL) and apolipoproteins for the binding and

CC in vivo transport of nucleic acids. The composition is used to deliver

CC nucleic acids to eukaryotic cells, in vivo or in vitro, for expressing

CC a therapeutic polypeptide or antisense molecule (or ribozyme).

CC Specifically they are used for gene therapy of cancers (particularly

CC non-small cell lung carcinoma), diabetes, cystic fibrosis and

CC arteriosclerosis.

QY Query Match 46.1%; Score 256; DB 20; Length 589;

Best Local Similarity 97.8%; Pred. No. 2.1e-43;

Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

QY 2 AGTTAATTAATAGTAAATCAATACGGGGGCTATTTAGTTCATAGCCCAATATGAGTTCCG 61

DB 588 AGTATTAATAGTATCAATTAACGGGGTCATTAGTTCATACCCCATATATGAGTTCCGC 529
QY 62 GTTACATTAACGTAAGTAAATGGCCCGGCTGACCGCCCAAGACCCCGCCCATG 121
DB 528 GTTACATTAACGTAAGTAAATGGCCCGGCTGACCGCCCAAGACCCCGCCCATG 469
QY 122 ACCTCAATTAATGACGTATGTTCCCATAGTAAGCCCAATAGGAGCTTCCATTGACGTCA 181
DB 468 ACCTCAATTAATGACGTATGTTCCCATAGTAAGCCCAATAGGAGCTTCCATTGACGTCA 409
QY 182 TGGGTGAGTATTTACGGTAACGTCCTCA-TTGGCAGTACATCAAGTGTATCATATGCCA 240
DB 408 TGGGTGAGTATTTACGGTAACGTCCTCA-TTGGCAGTACATCAAGTGTATCATATGCCA 349
QY 241 AGTACGCCCCCTATTGACGTCAATGACGTAATG 276
DB 348 AGTACGCCCCCTATTGACGTCAATGACGTAATG 313

RESULT 14

AAH24425 ID AAH24425 standard; DNA; 590 BP.

AAH24425;

02-AUG-2001 (first entry)

DE Cauliflower mosaic virus promoter.

KW Cauliflower mosaic virus; promoter; signal peptide; chloroella;
KW gene expression; protein production; human growth hormone; ds.

OS Cauliflower mosaic virus.

JP2000354490-A.

26-DEC-2000.

15-JUN-1999; 99JP-0168271.

15-JUN-1999; 99JP-0168271.

(TOYOTA) TOYOTA JIDOSHA KK.

WPI; 2001-275809/29.

New signal peptides useful for the preparation of human growth hormone
and transformed chloroella

Example 1; Page 12; 15pp; Japanese.

The present sequence is provided in a specification relating to signal
peptides for expression and secretion of a protein in chloroella. The
peptides are of the formula:Met-Ala-Asn-Lys-X₁-(Leu)_n-X₂-Ala-Ser-Gly.X₁ - Ser or Leu;X₂ - an integer of 5-15;X_{1,2} - Gly-Ser-Leu or Pro-Leu-Ala.The signal peptides are useful in the preparation of human growth
hormone and transformed chloroella. Signal peptides; DNA encoding the
peptides, gene expression cassettes, recombinant vectors containing the
cassettes, and transformants having the vectors are provided. The
present sequence is a promoter which may be used in the invention.

Sequence 590 BP; 149 A; 143 C; 141 G; 157 T; 0 other;

Query Match 46.1%; Score 256; DB 22; Length 590;

Best Local Similarity 97.8%; Pred. No. 2,3e-43;

Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

2 AGTATTAATAGTATCAATTAACGGGGTCATTAGTTCATACCCCATATATGAGTTCCGC 61

DB 2 AGTATTAATAGTATCAATTAACGGGGTCATTAGTTCATACCCCATATATGAGTTCCGC 61
QY 62 GTTACATTAACGTAAGTAAATGGCCCGGCTGACCGCCCAAGACCCCGCCCATG 121
DB 62 GTTACATTAACGTAAGTAAATGGCCCGGCTGACCGCCCAAGACCCCGCCCATG 121
QY 122 ACCTCAATTAATGACGTATGTTCCCATAGTAAGCCCAATAGGAGCTTCCATTGACGTCA 181
DB 122 ACCTCAATTAATGACGTATGTTCCCATAGTAAGCCCAATAGGAGCTTCCATTGACGTCA 181
QY 182 TGGGTGAGTATTTACGGTAACGTCCTCA-TTGGCAGTACATCAAGTGTATCATATGCCA 240
DB 182 TGGGTGAGTATTTACGGTAACGTCCTCA-TTGGCAGTACATCAAGTGTATCATATGCCA 241
QY 241 AGTACGCCCCCTATTGACGTCAATGACGTAATG 276
DB 242 AGTACGCCCCCTATTGACGTCAATGACGTAATG 277

RESULT 15

AA295643 ID AA295643 standard; DNA; 763 BP.

AA295643;

08-JUN-2000 (first entry)

DE Cytomegalovirus derived promoter sequence SEQ ID NO:1.

KW Cytomegalovirus; CMV; promoter; human growth hormone; PGH; ovine;

KW prostaglandin G/H synthase; plasmid; prostanoid; circular;

KW cardiant; thrombolytic; antiinflammatory; gene therapy;

KW pulmonary embolus; myocardial infarction; lung disease;

KW adult respiratory distress syndrome; ds.

Human cytomegalovirus.

US6030638-A.

29-FEB-2000.

02-JUN-1995; 95US-0459493.

19-AUG-1991; 91US-0746941.

21-JUN-1993; 93US-0080221.

(UYVA-) UNIV VANDERBILT.

Meyrick B, Canonico A, Brigham K, Conary JT;

WPI; 2000-205193/18.

Increasing prostanoid production in vivo comprising delivering and
PT hyperexpressing a prostaglandin synthase gene in cells; useful for
treating e.g. myocardial infarction and general lung disease -

Disclosure; Column 7-10; 11pp; English.

A method has been developed of increasing prostanoid production in vivo
comprising delivering and hyperexpressing a prostaglandin synthase gene
in cells. The method can be used for gene therapy. The method is useful
for the treatment of diseases such as pulmonary embolus, myocardial
infarction and general lung disease (e.g. adult respiratory distress
syndrome). Generally the present invention provides a plasmid comprising
a cytomegalovirus (CMV) derived promoter sequence driving the coding
region for ovine prostaglandin G/H synthase (PGH). In order to increase
the expression of the gene, the construct also contains a short
translation augmenting sequence and a portion of the 3' untranslated
region from the human growth hormone (hGH) gene. The present sequence
represents the CMV promoter sequence used in the exemplification of the
present invention.

Sequence 763 BP; 203 A; 174 C; 164 G; 222 T; 0 other;

Query Match 46.1%; Score 256; DB 21; Length 763;
 Best Local Similarity 97.8%; Pred. No. 2.1e-43;
 Matches 270; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

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OY 2 AGTTATTAATAGTATCAATTCACGGGGTCATTAGTTCATAGCCCATATATGAGTCCGC 61
    |||||||
DB 180 AGTTATTAATAGTATCAATTCACGGGGTCATTAGTTCATAGCCCATATATGAGTCCGC 239
    |||||||
OY 62 GTTACATAACTTACGTAATTTGGCCCGCGCTGACGCCACGACCCCGCCCATTTG 121
    |||||||
DB 240 GTTACATAACTTACGTAATTTGGCCCGCGCTGACGCCACGACCCCGCCCATTTG 299
    |||||||
OY 122 ACGTCAATAATAGCATATGTTCCCATAGTACGCCAATAGGACTTTCATTGACGTCAA 181
    |||||||
DB 300 ACGTCAATAATAGCATATGTTCCCATAGTACGCCAATAGGACTTTCATTGACGTCAA 359
    |||||||
OY 182 TGGGTGAGTATTTACGGTAACCTGCCA-TTGGCAGTACATCAAGTATCATATAGCCA 240
    |||||||
DB 360 TGGGTGAGTATTTACGGTAACCTGCCA-TTGGCAGTACATCAAGTATCATATAGCCA 419
    |||||||
OY 241 AGTACGCCCCCTATTGACGTCAATGACGGTAATG 276
    |||||||
DB 420 AGTACGCCCCCTATTGACGTCAATGACGGTAATG 455
    |||||||
    
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